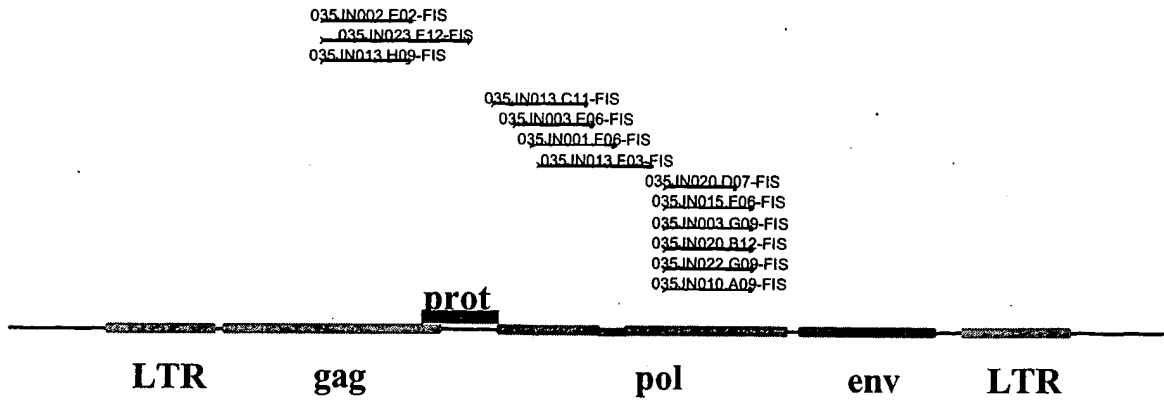
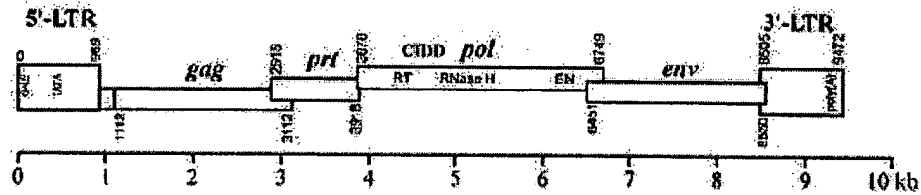
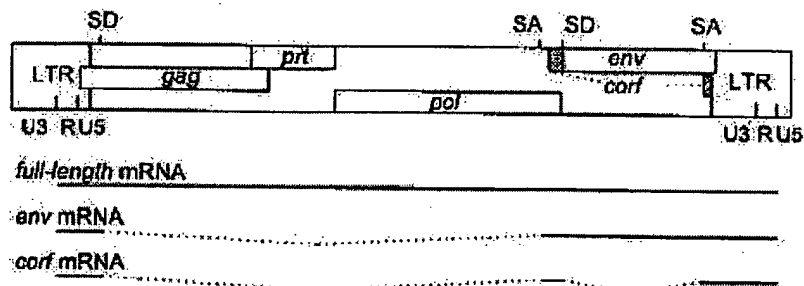


FIGURE 1**FIGURE 2****FIGURE 3**

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FIGURE 4

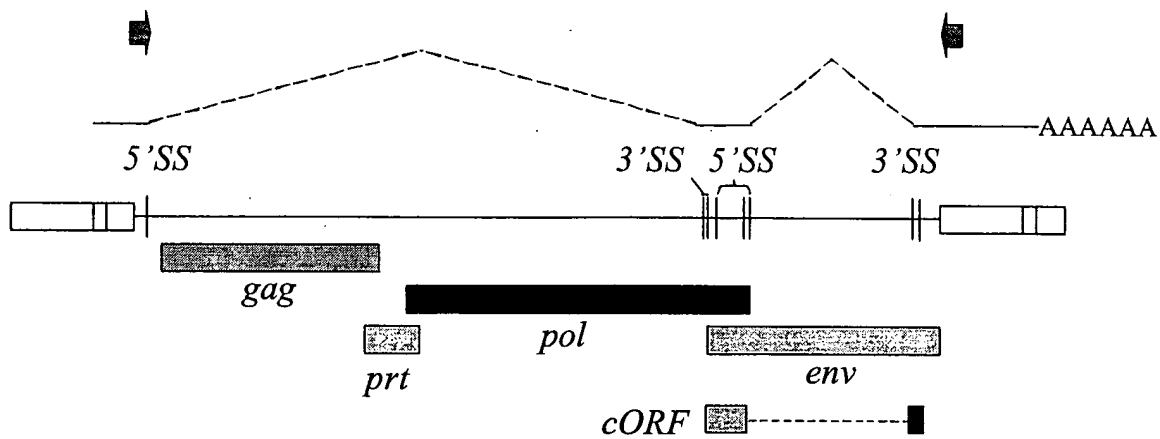
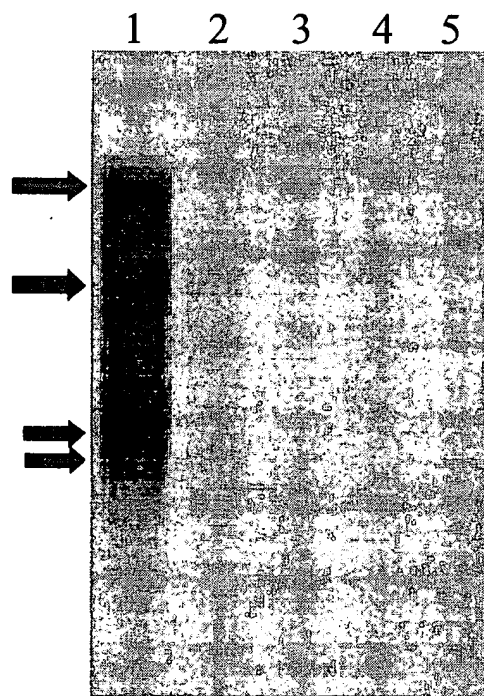


FIGURE 5



1004604-120704

ENV GENOMIC HERV MDA (1) -----

ENV GENOMIC HERV-K TAN. (1) -----

ENV GENOMIC AC025420 (1) -----

ENV GENOMIC AP000776 (1) -----

ENV GENOMIC HERV-K8 (1) -----

ENV GENOMIC HERV-KI (1) -----

ENV HERV-K AF023261 (1) GGGGAGAGGTTTGTGCTGTGTTTCACAGGAGAAAAATCAGCTTCTCGTTTGGATACCCATAGCACTTTGAGTTCTTAC

ENV GEN AL035086 (1) -----

ENV GENOMIC AL035587 (1) -----

ENV GENOMIC AC012068 (1) -----

ENV GENOMIC AF277315 (1) -----

ENV GENOMIC AF027650 (1) GGGGAGAGGTTTGTGCTGTGTTTCACAGGAGAAAAATCAGCTTCTCGTTTGGATACCCATAGCACTTTGAGTTCTTAC

ENV GENOMIC AC078899 (1) -----

ENV GENOMIC HERV-KII (1) -----

ENV GENOMIC AC008813 (1) -----

ENV GENOMIC AC012309 (1) -----

ENV GENOMIC AL121932 (1) -----

ENV GENOMIC AD000090 (1) -----

ENV GEN AL160008 (1) -----

ENV GENOMIC HEU32496 (1) ---GCGTAATCATTTGAGGACAAAGTCGACGAGAGAGATCCCGAGGAGCGTCTACAGTCAGCGCTTACACATTTGAGTTCTTAC

ENV GENOMIC AC011467 (1) ---GGTTTGTGCTGTGTTTCACAGGAGAGA-AAATCAGCTTCTCGTTTGGATACCCATAGCACTTTGAGTTCTTAC

ENV GENOMIC AF235103 (1) -----

ENV GENOMIC AC026786 (1) -----

ENV GENOMIC AC034203 (1) -----

ENV GENOMIC AC018809 (1) -----

ENV GENOMIC HERV-KI02 AF164610 (1) -----

ENV GENOMIC FRAG. AF260253 (1) -----

CONSENSUS (1) -----

ACATTTGAAGTTCTTAC

81

ENV GENOMIC HERV MDA (18) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC HERV-K TAN. (18) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AC025420 (18) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AP000776 (18) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC HERV-K8 (1) -----

ENV GENOMIC HERV-KI (18) ATGAACCTATCTGAGATGCAAGAA

ENV HERV-K AF023261 (81) ATGAACCTATCTGAGATGCAAGAA

ENV GEN AL035086 (1) -----

ENV GENOMIC AL035587 (18) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AC012068 (6) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AF277315 (15) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AF027650 (79) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AC078899 (16) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC HERV-KII (18) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AC008813 (29) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AC012309 (21) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AL121932 (18) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AD000090 (18) ATGAACCTATCTGAGATGCAAGAA

ENV GEN AL160008 (1) -----

ENV GENOMIC HEU32496 (78) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AC011467 (74) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AF235103 (72) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC AC026786 (58) -----

ENV GENOMIC AC034203 (58) -----

ENV GENOMIC AC018809 (16) -----

ENV GENOMIC HERV-KI02 AF164610 (70) ATGAACCTATCTGAGATGCAAGAA

ENV GENOMIC FRAG. AF260253 (1) -----

CONSENSUS (81) ATGAACCTATCTGAGATGCAAGAA

161

ENV GENOMIC HERV MDA (72) -----

ENV GENOMIC HERV-K TAN. (83) -----

ENV GENOMIC AC025420 (83) -----

ENV GENOMIC AP000776 (83) -----

ENV GENOMIC HERV-K8 (15) -----

ENV GENOMIC HERV-KI (83) -----

ENV HERV-K AF023261 (146) -----

ENV GEN AL035086 (1) -----

ENV GENOMIC AL035587 (83) -----

ENV GENOMIC AC012068 (71) -----

ENV GENOMIC AF277315 (80) -----

ENV GENOMIC AF027650 (144) -----

ENV GENOMIC AC078899 (81) -----

ENV GENOMIC HERV-KII (72) -----

ENV GENOMIC AC008813 (104) -----

ENV GENOMIC AC012309 (86) -----

ENV GENOMIC AL121932 (83) -----

ENV GENOMIC AD000090 (88) -----

ENV GEN AL160008 (1) -----

ENV GENOMIC HEU32496 (143) -----

ENV GENOMIC AC011467 (128) -----

ENV GENOMIC AF235103 (142) -----

ENV GENOMIC AC026786 (58) -----

ENV GENOMIC AC034203 (123) -----

ENV GENOMIC AC018809 (70) -----

ENV GENOMIC HERV-KI02 AF164610 (124) -----

ENV GENOMIC FRAG. AF260253 (1) -----

CONSENSUS (161) T GACTCACAAGATGAA AAAATGGTGA TCAGAAGAACAGATGAAGTTGCCATCCACCAAGAA GC GA

240

T GACTCACAAGATGAA AAAATGGTGA TCAGAAGAACAGATGAAGTTGCCATCCACCAAGAA GC GA

[illegible]

[illegible]

FIGURE 6 CONTD...

ENV GENOMIC HERV MDA	(609)	721	800
ENV GENOMIC HERV-K TAN.	(630)	---	---
ENV GENOMIC AC025420	(627)	---	---
ENV GENOMIC AP000776	(630)	---	---
ENV GENOMIC HERV-K8	(291)	---	---
ENV GENOMIC HERV-KI	(630)	---	---
ENV HERV-K AF023261	(693)	---	---
ENV GEN AL035086	(416)	---	---
ENV GENOMIC AL035587	(630)	---	---
ENV GENOMIC AC012068	(618)	---	---
ENV GENOMIC AF277315	(627)	---	---
ENV GENOMIC AF027650	(692)	---	---
ENV GENOMIC AC078899	(629)	---	---
ENV GENOMIC HERV-KII	(338)	---	---
ENV GENOMIC AC008813	(650)	---	---
ENV GENOMIC AC012309	(629)	---	---
ENV GENOMIC AL121932	(627)	---	---
ENV GENOMIC AD000090	(643)	---	---
ENV GEN AL160008	(482)	---	---
ENV GENOMIC HEU32496	(441)	---	---
ENV GENOMIC AC011467	(394)	---	---
ENV GENOMIC AF235103	(688)	---	---
ENV GENOMIC AC026786	(564)	---	---
ENV GENOMIC AC034203	(670)	---	---
ENV GENOMIC AC018809	(337)	---	---
ENV GENOMIC HERV-KI02 AF164610	(390)	---	---
ENV GENOMIC FRAG. AF260253	(1)	---	---
CONSENSUS	(721)	C GGTAAAT ATTACA GACTTTTCTATCAAGATCATTAATAATTAG CCTAAGGGGAACCTTGCCCAAGGAAT	
ENV GENOMIC HERV MDA	(685)	801	880
ENV GENOMIC HERV-K TAN.	(710)	---	---
ENV GENOMIC AC025420	(707)	---	---
ENV GENOMIC AP000776	(710)	---	---
ENV GENOMIC HERV-K8	(291)	---	---
ENV GENOMIC HERV-KI	(710)	---	---
ENV HERV-K AF023261	(701)	---	---
ENV GEN AL035086	(496)	---	---
ENV GENOMIC AL035587	(710)	---	---
ENV GENOMIC AC012068	(698)	---	---
ENV GENOMIC AF277315	(707)	---	---
ENV GENOMIC AF027650	(700)	---	---
ENV GENOMIC AC078899	(709)	---	---
ENV GENOMIC HERV-KII	(418)	---	---
ENV GENOMIC AC008813	(729)	---	---
ENV GENOMIC AC012309	(709)	---	---
ENV GENOMIC AL121932	(707)	---	---
ENV GENOMIC AD000090	(723)	---	---
ENV GEN AL160008	(543)	---	---
ENV GENOMIC HEU32496	(441)	---	---
ENV GENOMIC AC011467	(474)	---	---
ENV GENOMIC AF235103	(768)	---	---
ENV GENOMIC AC026786	(644)	---	---
ENV GENOMIC AC034203	(750)	---	---
ENV GENOMIC AC018809	(417)	---	---
ENV GENOMIC HERV-KI02 AF164610	(470)	---	---
ENV GENOMIC FRAG. AF260253	(1)	---	---
CONSENSUS	(801)	TCCTAAG ATCAAAA A CAGAAGTTTATGTTGGGAAGATGTGGC AATAGTGC GTGATATTACAAAACATG	
ENV GENOMIC HERV MDA	(764)	881	960
ENV GENOMIC HERV-K TAN.	(790)	---	---
ENV GENOMIC AC025420	(787)	---	---
ENV GENOMIC AP000776	(790)	---	---
ENV GENOMIC HERV-K8	(291)	---	---
ENV GENOMIC HERV-KI	(790)	---	---
ENV HERV-K AF023261	(701)	---	---
ENV GEN AL035086	(576)	---	---
ENV GENOMIC AL035587	(790)	---	---
ENV GENOMIC AC012068	(778)	---	---
ENV GENOMIC AF277315	(787)	---	---
ENV GENOMIC AF027650	(700)	---	---
ENV GENOMIC AC078899	(789)	---	---
ENV GENOMIC HERV-KII	(498)	---	---
ENV GENOMIC AC008813	(809)	---	---
ENV GENOMIC AC012309	(789)	---	---
ENV GENOMIC AL121932	(807)	---	---
ENV GENOMIC AD000090	(803)	---	---
ENV GEN AL160008	(623)	---	---
ENV GENOMIC HEU32496	(441)	---	---
ENV GENOMIC AC011467	(554)	---	---
ENV GENOMIC AF235103	(848)	---	---
ENV GENOMIC AC026786	(724)	---	---
ENV GENOMIC AC034203	(830)	---	---
ENV GENOMIC AC018809	(497)	---	---
ENV GENOMIC HERV-KI02 AF164610	(550)	---	---
ENV GENOMIC FRAG. AF260253	(1)	---	---
CONSENSUS	(881)	AATTGGAACTATTATAGATTGGGCACCTCGAGGTCATTTCTA CACA ATTGCA	

FIGURE 6 CONTD...

ENV GENOMIC HERV MDA	(820)	961	1040
ENV GENOMIC HERV-K TAN.	(846)	GG CAACTCA TC TGCC AG GCACAAG	AGTCCAGCTGTTGATAG GACTTAACAGAAAGT T
ENV GENOMIC AC025420	(843)		
ENV GENOMIC AP000776	(846)		
ENV GENOMIC HERV-K8	(291)		
ENV GENOMIC HERV-KI	(846)		
ENV HERV-K AF023261	(701)		
ENV GEN AL035086	(632)		
ENV GENOMIC AL035587	(870)		
ENV GENOMIC AC012068	(834)		
ENV GENOMIC AF277315	(843)		
ENV GENOMIC AF027650	(700)		
ENV GENOMIC AC078899	(845)		
ENV GENOMIC HERV-KII	(554)		
ENV GENOMIC AC008813	(865)		
ENV GENOMIC AC012309	(845)		
ENV GENOMIC AL121932	(843)		
ENV GENOMIC AD000090	(859)		
ENV GEN AL160008	(647)		
ENV GENOMIC HEU32496	(441)		
ENV GENOMIC AC011467	(610)		
ENV GENOMIC AF235103	(904)		
ENV GENOMIC AC026786	(780)		
ENV GENOMIC AC034203	(886)		
ENV GENOMIC AC018809	(553)		
ENV GENOMIC HERV-KI02	AF164610		
ENV GENOMIC FRAG.	AF260253		
CONSENSUS	(961)		
ENV GENOMIC HERV MDA	(900)	1041	1120
ENV GENOMIC HERV-K TAN.	(914)	AGAG AA T A TA AA TTA A TC TCTA CC TGG AATGGG GAAAA GGAAT TC C	
ENV GENOMIC AC025420	(911)		
ENV GENOMIC AP000776	(914)		
ENV GENOMIC HERV-K8	(291)		
ENV GENOMIC HERV-KI	(914)		
ENV HERV-K AF023261	(701)		
ENV GEN AL035086	(700)		
ENV GENOMIC AL035587	(950)		
ENV GENOMIC AC012068	(914)		
ENV GENOMIC AF277315	(923)		
ENV GENOMIC AF027650	(700)		
ENV GENOMIC AC078899	(913)		
ENV GENOMIC HERV-KII	(622)		
ENV GENOMIC AC008813	(933)		
ENV GENOMIC AC012309	(913)		
ENV GENOMIC AL121932	(911)		
ENV GENOMIC AD000090	(927)		
ENV GEN AL160008	(647)		
ENV GENOMIC HEU32496	(441)		
ENV GENOMIC AC011467	(676)		
ENV GENOMIC AF235103	(984)		
ENV GENOMIC AC026786	(860)		
ENV GENOMIC AC034203	(966)		
ENV GENOMIC AC018809	(621)		
ENV GENOMIC HERV-KI02	AF164610		
ENV GENOMIC FRAG.	AF260253		
CONSENSUS	(1041)		
ENV GENOMIC HERV MDA	(967)	1121	1200
ENV GENOMIC HERV-K TAN.	(984)		
ENV GENOMIC AC025420	(981)		
ENV GENOMIC AP000776	(984)		
ENV GENOMIC HERV-K8	(291)		
ENV GENOMIC HERV-KI	(984)		
ENV HERV-K AF023261	(701)		
ENV GEN AL035086	(770)		
ENV GENOMIC AL035587	(1017)		
ENV GENOMIC AC012068	(981)		
ENV GENOMIC AF277315	(990)		
ENV GENOMIC AF027650	(700)		
ENV GENOMIC AC078899	(993)		
ENV GENOMIC HERV-KII	(692)		
ENV GENOMIC AC008813	(1003)		
ENV GENOMIC AC012309	(983)		
ENV GENOMIC AL121932	(981)		
ENV GENOMIC AD000090	(997)		
ENV GEN AL160008	(647)		
ENV GENOMIC HEU32496	(441)		
ENV GENOMIC AC011467	(683)		
ENV GENOMIC AF235103	(1051)		
ENV GENOMIC AC026786	(927)		
ENV GENOMIC AC034203	(1033)		
ENV GENOMIC AC018809	(691)		
ENV GENOMIC HERV-KI02	AF164610		
ENV GENOMIC FRAG.	AF260253		
CONSENSUS	(1121)		

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FIGURE 6 CONTD...

	1201		1280
ENV GENOMIC HERV MDA	(967)	-----	-----
ENV GENOMIC HERV-K TAN.	(984)	-----	-----
ENV GENOMIC AC025420	(981)	-----	-----
ENV GENOMIC AP000776	(984)	-----	-----
ENV GENOMIC HERV-K8	(291)	-----	-----
ENV GENOMIC HERV-KI	(984)	-----	-----
ENV HERV-K AF023261	(701)	-----	-----
ENV GEN AL035086	(770)	-----	-----
ENV GENOMIC AL035587	(1017)	-----	-----
ENV GENOMIC AC012068	(981)	-----	-----
ENV GENOMIC AF277315	(990)	-----	-----
ENV GENOMIC AF027650	(700)	-----	-----
ENV GENOMIC AC078899	(1073)	CCACCCCGACTAACGCACATGCCCACTAGGGCGTGTCACTCAGAAGTGTGAAACTCAACCGATCCCGCCCTACCCCG	-----
ENV GENOMIC HERV-KII	(692)	-----	-----
ENV GENOMIC AC008813	(1003)	-----	-----
ENV GENOMIC AC012309	(983)	-----	-----
ENV GENOMIC AL121932	(981)	-----	-----
ENV GENOMIC AD000090	(997)	-----	-----
ENV GEN AL160008	(647)	-----	-----
ENV GENOMIC HEU32496	(441)	-----	-----
ENV GENOMIC AC011467	(683)	-----	-----
ENV GENOMIC AF235103	(1051)	-----	-----
ENV GENOMIC AC026786	(927)	-----	-----
ENV GENOMIC AC034203	(1033)	-----	-----
ENV GENOMIC AC018809	(691)	-----	-----
ENV GENOMIC HERV-KI02 AF164610	(744)	-----	-----
ENV GENOMIC FRAG. AF260253	(1)	-----	-----
CONSENSUS	(1201)	-----	-----
	1281		1360
ENV GENOMIC HERV MDA	(967)	-----	-----
ENV GENOMIC HERV-K TAN.	(984)	-----	-----
ENV GENOMIC AC025420	(981)	-----	-----
ENV GENOMIC AP000776	(984)	-----	-----
ENV GENOMIC HERV-K8	(291)	-----	-----
ENV GENOMIC HERV-KI	(984)	-----	-----
ENV HERV-K AF023261	(701)	-----	-----
ENV GEN AL035086	(770)	-----	-----
ENV GENOMIC AL035587	(1017)	-----	-----
ENV GENOMIC AC012068	(981)	-----	-----
ENV GENOMIC AF277315	(990)	-----	-----
ENV GENOMIC AF027650	(700)	-----	-----
ENV GENOMIC AC078899	(1153)	ACCACTCCTCACCCAGCATCCATAAAAGCGCGCTGCACCTTTGCGCACAGCGTGACTTCCCTGGCGGACCAGTGAACCTC	-----
ENV GENOMIC HERV-KII	(692)	-----	-----
ENV GENOMIC AC008813	(1003)	-----	-----
ENV GENOMIC AC012309	(983)	-----	-----
ENV GENOMIC AL121932	(981)	-----	-----
ENV GENOMIC AD000090	(997)	-----	-----
ENV GEN AL160008	(647)	-----	-----
ENV GENOMIC HEU32496	(441)	-----	-----
ENV GENOMIC AC011467	(683)	-----	-----
ENV GENOMIC AF235103	(1051)	-----	-----
ENV GENOMIC AC026786	(927)	-----	-----
ENV GENOMIC AC034203	(1033)	-----	-----
ENV GENOMIC AC018809	(691)	-----	-----
ENV GENOMIC HERV-KI02 AF164610	(744)	-----	-----
ENV GENOMIC FRAG. AF260253	(1)	-----	-----
CONSENSUS	(1281)	-----	-----
	1361		1440
ENV GENOMIC HERV MDA	(967)	-----	-----
ENV GENOMIC HERV-K TAN.	(984)	-----	-----
ENV GENOMIC AC025420	(981)	-----	-----
ENV GENOMIC AP000776	(984)	-----	-----
ENV GENOMIC HERV-K8	(291)	-----	-----
ENV GENOMIC HERV-KI	(984)	-----	-----
ENV HERV-K AF023261	(701)	-----	-----
ENV GEN AL035086	(770)	-----	-----
ENV GENOMIC AL035587	(1017)	-----	-----
ENV GENOMIC AC012068	(981)	-----	-----
ENV GENOMIC AF277315	(990)	-----	-----
ENV GENOMIC AF027650	(700)	-----	-----
ENV GENOMIC AC078899	(1233)	ACCGGAGAGCTCAATAAAGAAAGATTTTTGCCCTCTTTGTCTTGCCCTTTGGCCTTATTGATCCACGGTGCCCTTTCCATTG	-----
ENV GENOMIC HERV-KII	(692)	-----	-----
ENV GENOMIC AC008813	(1003)	-----	-----
ENV GENOMIC AC012309	(983)	-----	-----
ENV GENOMIC AL121932	(981)	-----	-----
ENV GENOMIC AD000090	(997)	-----	-----
ENV GEN AL160008	(647)	-----	-----
ENV GENOMIC HEU32496	(441)	-----	-----
ENV GENOMIC AC011467	(683)	-----	-----
ENV GENOMIC AF235103	(1051)	-----	-----
ENV GENOMIC AC026786	(927)	-----	-----
ENV GENOMIC AC034203	(1033)	-----	-----
ENV GENOMIC AC018809	(691)	-----	-----
ENV GENOMIC HERV-KI02 AF164610	(744)	-----	-----
ENV GENOMIC FRAG. AF260253	(1)	-----	-----
CONSENSUS	(1361)	-----	-----

FIGURE 6 contd...

ENV GENOMIC HERV MDA	(967)	1441	1520
ENV GENOMIC HERV-K TAN.	(984)	-----	-----
ENV GENOMIC AC025420	(981)	-----	-----
ENV GENOMIC AP000776	(984)	-----	-----
ENV GENOMIC HERV-K8	(291)	-----	-----
ENV GENOMIC HERV-KI	(984)	-----	-----
ENV HERV-K AF023261	(701)	-----	-----
ENV GEN AL035086	(770)	-----	-----
ENV GENOMIC AL035587	(1017)	-----	-----
ENV GENOMIC AC012068	(981)	-----	-----
ENV GENOMIC AF277315	(990)	-----	-----
ENV GENOMIC AF027650	(700)	-----	-----
ENV GENOMIC AC078899	(1313)	-----	-----
ENV GENOMIC HERV-KII	(692)	-----	-----
ENV GENOMIC AC008813	(1003)	-----	-----
ENV GENOMIC AC012309	(983)	-----	-----
ENV GENOMIC AL121932	(981)	-----	-----
ENV GENOMIC AD000090	(997)	-----	-----
ENV GEN AL160008	(647)	-----	-----
ENV GENOMIC HEU32496	(441)	-----	-----
ENV GENOMIC AC011467	(683)	-----	-----
ENV GENOMIC AF235103	(1051)	-----	-----
ENV GENOMIC AC026786	(927)	-----	-----
ENV GENOMIC AC034203	(1033)	-----	-----
ENV GENOMIC AC018809	(691)	-----	-----
ENV GENOMIC HERV-K102 AF164610	(744)	-----	-----
ENV GENOMIC FRAG. AF260253	(1)	-----	-----
CONSENSUS	(1441)	CC GACCAAA TA T AGTCTGTT CTGGTCTGAACATCCAGAATTATGGA GCTTACTGTGGCC	
ENV GENOMIC HERV MDA	(1031)	1521	1600
ENV GENOMIC HERV-K TAN.	(1053)	-----	-----
ENV GENOMIC AC025420	(1050)	-----	-----
ENV GENOMIC AP000776	(1053)	-----	-----
ENV GENOMIC HERV-K8	(291)	-----	-----
ENV GENOMIC HERV-KI	(1053)	-----	-----
ENV HERV-K AF023261	(701)	-----	-----
ENV GEN AL035086	(839)	-----	-----
ENV GENOMIC AL035587	(1086)	-----	-----
ENV GENOMIC AC012068	(1046)	-----	-----
ENV GENOMIC AF277315	(1059)	-----	-----
ENV GENOMIC AF027650	(700)	-----	-----
ENV GENOMIC AC078899	(1393)	-----	-----
ENV GENOMIC HERV-KII	(757)	-----	-----
ENV GENOMIC AC008813	(1072)	-----	-----
ENV GENOMIC AC012309	(1052)	-----	-----
ENV GENOMIC AL121932	(1050)	-----	-----
ENV GENOMIC AD000090	(1066)	-----	-----
ENV GEN AL160008	(647)	-----	-----
ENV GENOMIC HEU32496	(441)	-----	-----
ENV GENOMIC AC011467	(727)	-----	-----
ENV GENOMIC AF235103	(1120)	-----	-----
ENV GENOMIC AC026786	(996)	-----	-----
ENV GENOMIC AC034203	(1102)	-----	-----
ENV GENOMIC AC018809	(760)	-----	-----
ENV GENOMIC HERV-K102 AF164610	(813)	-----	-----
ENV GENOMIC FRAG. AF260253	(1)	-----	-----
CONSENSUS	(1521)	TCA ACCACATTAGAAATTTGGTCTGGAATCAA CT TAGAACAAGAGATC TAAGCCAT TTATACTATC ACCTAA	
ENV GENOMIC HERV MDA	(1110)	1601	1680
ENV GENOMIC HERV-K TAN.	(1132)	-----	-----
ENV GENOMIC AC025420	(1129)	-----	-----
ENV GENOMIC AP000776	(1132)	-----	-----
ENV GENOMIC HERV-K8	(291)	-----	-----
ENV GENOMIC HERV-KI	(1132)	-----	-----
ENV HERV-K AF023261	(701)	-----	-----
ENV GEN AL035086	(918)	-----	-----
ENV GENOMIC AL035587	(1165)	-----	-----
ENV GENOMIC AC012068	(1125)	-----	-----
ENV GENOMIC AF277315	(1138)	-----	-----
ENV GENOMIC AF027650	(700)	-----	-----
ENV GENOMIC AC078899	(1472)	-----	-----
ENV GENOMIC HERV-KII	(836)	-----	-----
ENV GENOMIC AC008813	(1151)	-----	-----
ENV GENOMIC AC012309	(1131)	-----	-----
ENV GENOMIC AL121932	(1130)	-----	-----
ENV GENOMIC AD000090	(1143)	-----	-----
ENV GEN AL160008	(647)	-----	-----
ENV GENOMIC HEU32496	(441)	-----	-----
ENV GENOMIC AC011467	(806)	-----	-----
ENV GENOMIC AF235103	(1199)	-----	-----
ENV GENOMIC AC026786	(1075)	-----	-----
ENV GENOMIC AC034203	(1181)	-----	-----
ENV GENOMIC AC018809	(839)	-----	-----
ENV GENOMIC HERV-K102 AF164610	(892)	-----	-----
ENV GENOMIC FRAG. AF260253	(1)	-----	-----
CONSENSUS	(1601)	ATTCCAGTCT ACA TTCTTT CAAGTTTG GTAAGGCCCTTATAT GCTAGTTGTAGGAATA TAGTATTAA	

ENV GENOMIC HERV MDA (1190) 1761
 ENV GENOMIC HERV-K TAN. (1209) 1760
 ENV GENOMIC AC025420 (1206)
 ENV GENOMIC AP000776 (1209)
 ENV GENOMIC HERV-K8 (291)
 ENV GENOMIC HERV-KI (1209)
 ENV HERV-K AF023261 (701)
 ENV GEN AL035086 (995)
 ENV GENOMIC AL035587 (1242)
 ENV GENOMIC AC012068 (1202)
 ENV GENOMIC AF277315 (1215)
 ENV GENOMIC AF027650 (700)
 ENV GENOMIC AC078899 (1549)
 ENV GENOMIC HERV-KII (913)
 ENV GENOMIC AC008813 (1228)
 ENV GENOMIC AC012309 (1208)
 ENV GENOMIC AL121932 (1207)
 ENV GENOMIC AD000090 (1220)
 ENV GEN AL160008 (647)
 ENV GENOMIC HERV32496 (441)
 ENV GENOMIC AC011467 (883)
 ENV GENOMIC AF235103 (1276)
 ENV GENOMIC AC026786 (1152)
 ENV GENOMIC AC034203 (1258)
 ENV GENOMIC AC018809 (916)
 ENV GENOMIC HERV-KI02 AF164610 (969)
 ENV GENOMIC FRAG. AF260253 (1)
 CONSENSUS (1681)
 CAGA TCCCAAACTATA ACCTGTGAAAATGTAGATTGTTTACTTGCATTGATTCAACTTTTAAATTGGCAGCAGCGT

ENV GENOMIC HERV MDA (1268) 1761
 ENV GENOMIC HERV-K TAN. (1287) 1840
 ENV GENOMIC AC025420 (1284)
 ENV GENOMIC AP000776 (1287)
 ENV GENOMIC HERV-K8 (291)
 ENV GENOMIC HERV-KI (1287)
 ENV HERV-K AF023261 (701)
 ENV GEN AL035086 (1073)
 ENV GENOMIC AL035587 (1320)
 ENV GENOMIC AC012068 (1280)
 ENV GENOMIC AF277315 (1293)
 ENV GENOMIC AF027650 (700)
 ENV GENOMIC AC078899 (1627)
 ENV GENOMIC HERV-KII (991)
 ENV GENOMIC AC008813 (1238)
 ENV GENOMIC AC012309 (1288)
 ENV GENOMIC AL121932 (1285)
 ENV GENOMIC AD000090 (1298)
 ENV GEN AL160008 (647)
 ENV GENOMIC HERV32496 (441)
 ENV GENOMIC AC011467 (961)
 ENV GENOMIC AF235103 (1354)
 ENV GENOMIC AC026786 (1230)
 ENV GENOMIC AC034203 (1336)
 ENV GENOMIC AC018809 (992)
 ENV GENOMIC HERV-KI02 AF164610 (1047)
 ENV GENOMIC FRAG. AF260253 (1)
 CONSENSUS (1761)
 ATTCTGCT GTGAGAGCAAGAGA GG GTGTGATCCCTGTGTCCATGACGACCGCTGGAGGC TC CCATCC TCCA

ENV GENOMIC HERV MDA (1348) 1841
 ENV GENOMIC HERV-K TAN. (1367) 1920
 ENV GENOMIC AC025420 (1364)
 ENV GENOMIC AP000776 (1367)
 ENV GENOMIC HERV-K8 (291)
 ENV GENOMIC HERV-KI (1367)
 ENV HERV-K AF023261 (701)
 ENV GEN AL035086 (1153)
 ENV GENOMIC AL035587 (1399)
 ENV GENOMIC AC012068 (1360)
 ENV GENOMIC AF277315 (1373)
 ENV GENOMIC AF027650 (700)
 ENV GENOMIC AC078899 (1707)
 ENV GENOMIC HERV-KII (1071)
 ENV GENOMIC AC008813 (1238)
 ENV GENOMIC AC012309 (1368)
 ENV GENOMIC AL121932 (1365)
 ENV GENOMIC AD000090 (1378)
 ENV GEN AL160008 (647)
 ENV GENOMIC HERV32496 (441)
 ENV GENOMIC AC011467 (1041)
 ENV GENOMIC AF235103 (1434)
 ENV GENOMIC AC026786 (1310)
 ENV GENOMIC AC034203 (1403)
 ENV GENOMIC AC018809 (1072)
 ENV GENOMIC HERV-KI02 AF164610 (1127)
 ENV GENOMIC FRAG. AF260253 (1)
 CONSENSUS (1841)
 TATTATT AC GAAGTATTAAGAAG TT TAA TAGATCCAAAAGATTCAATTTTACTTTAAATTCAGTGAATATGGG

FIGURE 6 CONTD...

ENV GENOMIC HERV MDA	(1428)	1921	2000
ENV GENOMIC HERV-K TAN.	(1447)		
ENV GENOMIC AC025420	(1444)		
ENV GENOMIC AP000776	(1447)		
ENV GENOMIC HERV-K8	(291)		
ENV GENOMIC HERV-KI	(1447)		
ENV HERV-K AF023261	(701)		
ENV GEN AL035086	(1233)		
ENV GENOMIC AL035587	(1475)		
ENV GENOMIC AC012068	(1440)		
ENV GENOMIC AF277315	(1453)		
ENV GENOMIC AF027650	(700)		
ENV GENOMIC AC078899	(1786)		
ENV GENOMIC HERV-KII	(1151)		
ENV GENOMIC AC008813	(1238)		
ENV GENOMIC AC012309	(1448)		
ENV GENOMIC AL121932	(1444)		
ENV GENOMIC AD000090	(1458)		
ENV GEN AL160008	(647)		
ENV GENOMIC HEU32496	(441)		
ENV GENOMIC AC011467	(1121)		
ENV GENOMIC AF235103	(1514)		
ENV GENOMIC AC026786	(1390)		
ENV GENOMIC AC034203	(1403)		
ENV GENOMIC AC018809	(1151)		
ENV GENOMIC HERV-K102 AF164610	(1207)		
ENV GENOMIC FRAG. AF260253	(1)		
CONSENSUS	(1921)	T ATTGCACTCACAGCTAC GCTCG G GC GGA TTGC TT CACTC TCTGTTCA C G A A T TGT AAT AT	
ENV GENOMIC HERV MDA	(1508)	2001	2080
ENV GENOMIC HERV-K TAN.	(1527)		
ENV GENOMIC AC025420	(1524)		
ENV GENOMIC AP000776	(1527)		
ENV GENOMIC HERV-K8	(291)		
ENV GENOMIC HERV-KI	(1527)		
ENV HERV-K AF023261	(701)		
ENV GEN AL035086	(1313)		
ENV GENOMIC AL035587	(1555)		
ENV GENOMIC AC012068	(1520)		
ENV GENOMIC AF277315	(1533)		
ENV GENOMIC AF027650	(700)		
ENV GENOMIC AC078899	(1788)		
ENV GENOMIC HERV-KII	(1231)		
ENV GENOMIC AC008813	(1238)		
ENV GENOMIC AC012309	(1528)		
ENV GENOMIC AL121932	(1524)		
ENV GENOMIC AD000090	(1538)		
ENV GEN AL160008	(647)		
ENV GENOMIC HEU32496	(441)		
ENV GENOMIC AC011467	(1201)		
ENV GENOMIC AF235103	(1594)		
ENV GENOMIC AC026786	(1470)		
ENV GENOMIC AC034203	(1403)		
ENV GENOMIC AC018809	(1231)		
ENV GENOMIC HERV-K102 AF164610	(1287)		
ENV GENOMIC FRAG. AF260253	(1)		
CONSENSUS	(2001)	TGCAAAA AA TTC CAA ATTGTGGAATTC CA A C AT GATCAAAAATTGGCAAAATCAAATTAATGATCTT	
ENV GENOMIC HERV MDA	(1586)	2081	2160
ENV GENOMIC HERV-K TAN.	(1605)		
ENV GENOMIC AC025420	(1602)		
ENV GENOMIC AP000776	(1605)		
ENV GENOMIC HERV-K8	(291)		
ENV GENOMIC HERV-KI	(1605)		
ENV HERV-K AF023261	(701)		
ENV GEN AL035086	(1392)		
ENV GENOMIC AL035587	(1633)		
ENV GENOMIC AC012068	(1598)		
ENV GENOMIC AF277315	(1611)		
ENV GENOMIC AF027650	(700)		
ENV GENOMIC AC078899	(1864)		
ENV GENOMIC HERV-KII	(1309)		
ENV GENOMIC AC008813	(1238)		
ENV GENOMIC AC012309	(1606)		
ENV GENOMIC AL121932	(1538)		
ENV GENOMIC AD000090	(1616)		
ENV GEN AL160008	(647)		
ENV GENOMIC HEU32496	(441)		
ENV GENOMIC AC011467	(1279)		
ENV GENOMIC AF235103	(1672)		
ENV GENOMIC AC026786	(1548)		
ENV GENOMIC AC034203	(1403)		
ENV GENOMIC AC018809	(1309)		
ENV GENOMIC HERV-K102 AF164610	(1365)		
ENV GENOMIC FRAG. AF260253	(1)		
CONSENSUS	(2081)	AGACAACTGTCATTGGATGGGAGA AG CTCATGAGCTT GAA ATC TTT CAGTTACA TGTGACTGGAATACGTC	

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FIGURE 6 CONTD...

ENV GENOMIC HERV MDA	(1664)	AGATTTTTGTATTACACCCAAAGCCATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT	2161	2240
ENV GENOMIC HERV-K TAN.	(1685)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AC025420	(1682)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AP000776	(1685)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC HERV-K8	(291)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC HERV-KI	(1685)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV HERV-K AF023261	(701)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GEN AL035086	(1472)	AGATTTTTGTATTACACCCAAAGCCATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AL035587	(1713)	AGATTTTTGTATTACACCCAAAGCCATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AC012068	(1678)	AGATTTTTGTATTACACCCAAAGCCATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AF277315	(1691)	AGATTTTTGTATTACACCCAAAGCCATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AF027650	(700)	AGATTTTTGTATTACACCCAAAGCCATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AC078899	(1944)	AGATTTTTGTATTACACCCAAAGCCATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC HERV-KII	(1389)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AC008813	(1238)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AC012309	(1682)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AL121932	(1538)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AD000090	(1696)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GEN AL160008	(647)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC HEU32496	(441)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AC011467	(1359)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AF235103	(1752)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AC026786	(1626)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AC034203	(1403)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC AC018809	(1389)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC HERV-KI02	AF164610	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC FRAG.	AF260253	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
CONSENSUS	(2161)	AGATTTTTGTATTACACCCAAATTTATAATGAGTCTGAGCATCACTGGGACATGGTTAGAT		
ENV GENOMIC HERV MDA	(1744)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA	2241	2320
ENV GENOMIC HERV-K TAN.	(1765)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AC025420	(1762)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AP000776	(1765)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC HERV-K8	(291)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC HERV-KI	(1765)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV HERV-K AF023261	(701)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GEN AL035086	(1552)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AL035587	(1793)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AC012068	(1758)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AF277315	(1771)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AF027650	(700)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AC078899	(2024)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC HERV-KII	(1469)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AC008813	(1238)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AC012309	(1762)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AL121932	(1538)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AD000090	(1776)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GEN AL160008	(647)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC HEU32496	(441)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AC011467	(1439)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AF235103	(1832)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AC026786	(1706)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AC034203	(1403)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC AC018809	(1468)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC HERV-KI02	AF164610	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC FRAG.	AF260253	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
CONSENSUS	(2241)	AGATAATCTCTGCTTAGACATTTTCAAATTAAGAGAA		
ENV GENOMIC HERV MDA	(1824)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG	2321	2400
ENV GENOMIC HERV-K TAN.	(1831)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AC025420	(1828)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AP000776	(1831)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC HERV-K8	(291)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC HERV-KI	(1831)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV HERV-K AF023261	(701)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GEN AL035086	(1618)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AL035587	(1859)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AC012068	(1824)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AF277315	(1837)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AF027650	(700)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AC078899	(2090)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC HERV-KII	(1535)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AC008813	(1238)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AC012309	(1826)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AL121932	(1538)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AD000090	(1842)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GEN AL160008	(647)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC HEU32496	(441)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AC011467	(1505)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AF235103	(1898)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AC026786	(1772)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AC034203	(1403)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC AC018809	(1534)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC HERV-KI02	AF164610	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
ENV GENOMIC FRAG.	AF260253	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		
CONSENSUS	(2321)	TAATTTGGTGGCAGGAACCTGAGGCAATTCAGGATTTGGCTGATGGCTTGAATCTTAACCTGTGGTTAG		

ENV GENOMIC HERV MDA (1904) AGCATCAGAAGTTCATGTTTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC HERV-K TAN. (1911) ACCATTCGAAGTATGTAAGATTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AC025420 (1908) ACCATTCGAAGTATGTAAGATTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AP000776 (1911) ACCATTCGAAGTATGTAAGATTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC HERV-K8 (291) -----
ENV GENOMIC HERV-KI (1906) ACCATTCGAAGTATGTAAGATTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV HERV-K AF023261 (701) -----
ENV GEN AL035086 (1698) AGCATCAGAAGTTCATGTTTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AL035587 (1933) AGCATCAGAAGTTCATGTTTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AC012068 (1904) ACCATTCGAAGTATGTAAGATTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AF277315 (1917) AGCATCAGAAGTTCATGTTTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AF027650 (700) -----
ENV GENOMIC AC078899 (2170) ACCATTCGAAGTATGTAAGATTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC HERV-KII (1615) ACCATTCGAAGTATGTAAGATTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AC008813 (1238) -----
ENV GENOMIC AC012309 (1905) ACCATTCGAAGTATGTAAGATTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AL121932 (1538) -----
ENV GENOMIC AD000090 (1922) ACCATTCGAAGTATGTAAGATTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GEN AL160008 (647) -----
ENV GENOMIC HEU32496 (441) -----
ENV GENOMIC AC011467 (1585) AGCATCAGAAGTTCATGTTTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AF235103 (1978) AGCATCAGAAGTTCATGTTTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AC026786 (1852) AGCATCAGAAGTTCATGTTTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC AC034203 (1403) -----
ENV GENOMIC AC018809 (1614) AGCATCAGAAGTTCATGTTTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC HERV-KI02 AF164610 (1671) AGCATCAGAAGTTCATGTTTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
ENV GENOMIC FRAG. AF260253 (151) AGCATCAGAAGTTCATGTTTAAATTCATATTAAATCCCTGTCGCTGTTTGT AGTCTCAGAGTGT
CONSENSUS (2401) ACCAT GAAGT C AC ATT TAAAT TCATATTAAATCCCTGT TGCTGTGT TGCTGTGTGT AGTCT CAGGTGT

2481 2560

ENV GENOMIC HERV MDA (1982) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC HERV-K TAN. (1989) ACCGAAACAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AC025420 (1886) ACCGAAACAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AP000776 (1989) ACCGAAACAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC HERV-K8 (291) -----
ENV GENOMIC HERV-KI (1984) ACCGAAACAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV HERV-K AF023261 (701) -----
ENV GEN AL035086 (1776) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AL035587 (2011) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AC012068 (1982) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AF277315 (1995) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AF027650 (700) -----
ENV GENOMIC AC078899 (2248) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC HERV-KII (1693) ACCGAAACAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AC008813 (1238) -----
ENV GENOMIC AC012309 (1983) ACCGAAACAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AL121932 (1538) -----
ENV GENOMIC AD000090 (2000) ACCGAAACAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GEN AL160008 (647) -----
ENV GENOMIC HEU32496 (441) -----
ENV GENOMIC AC011467 (1663) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AF235103 (2056) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AC026786 (1930) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC AC034203 (1403) -----
ENV GENOMIC AC018809 (1694) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC HERV-KI02 AF164610 (1749) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
ENV GENOMIC FRAG. AF260253 (228) ATCTCGCAGCTCCGAGAGACAGGACCGAGAGAAATGGGCCATAGTCGAGTGTGTTTGTGCAAAAAGAAAGGGGG
CONSENSUS (2481) A CCA CAGCTCCGAGAGACAGGACCA C AGAAGCGGCATGATGAGTGTGTTTGTGCAAAAAGAAAGGGGG

2561 264

ENV GENOMIC HERV MDA (2062) GGTATGTAAGAAAG --- AGAGATCAGTTCCTACTG STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC HERV-K TAN. (2069) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC AC025420 (2066) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC AP000776 (2069) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC HERV-K8 (291) -----
ENV GENOMIC HERV-KI (2064) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV HERV-K AF023261 (701) -----
ENV GEN AL035086 (1856) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC AL035587 (2086) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC AC012068 (2062) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC AF277315 (2075) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC AF027650 (700) -----
ENV GENOMIC AC078899 (2328) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC HERV-KII (1773) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC AC008813 (1238) -----
ENV GENOMIC AC012309 (2063) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC AL121932 (1538) -----
ENV GENOMIC AD000090 (2080) AATATG --- GGAAGAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GEN AL160008 (647) -----
ENV GENOMIC HEU32496 (441) -----
ENV GENOMIC AC011467 (1699) -----
ENV GENOMIC AF235103 (2136) --TTATGTAGGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC AC026786 (2010) --TTATGTAGGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC AC034203 (1403) -----
ENV GENOMIC AC018809 (1774) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC HERV-KI02 AF164610 (1829) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
ENV GENOMIC FRAG. AF260253 (308) --AAAGTCCGAAAGACAGAGAGATCAGTTGTACTGT STCTAGTAGAGT AGAGACATTAAGACTCCATT
CONSENSUS (2561) A ATGT GGAAGAG AGAGAGATCAG TTTACTGT GTCT TOTAGAAA A G AGACATTA GAGACTCCATT

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FIGURE 6 contd...

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ENV GENOMIC HERV MDA (2136) AGATCGTACTAA-----

ENV GENOMIC HERV-K TAN. (2146) GTTTAGTACTAA-----

ENV GENOMIC AC025420 (2143) GTTTAGTACTAA-----

ENV GENOMIC AP000776 (2146) GTTTAGTACTAA-----

ENV GENOMIC HERV-K8 (291) -----

ENV GENOMIC HERV-KI (2141) GTTTAGTACTAA-----

ENV HERV-K AF023261 (701) -----

ENV GEN AL035086 (1931) AGATCGTACTAA-----

ENV GENOMIC AL035587 (2146) -----

ENV GENOMIC AC012068 (2138) GAAAAAGAGCTGTACTTTGAACAATT-----

ENV GENOMIC AF277315 (2152) AGACCGT-----

ENV GENOMIC AF027650 (700) -----

ENV GENOMIC AC078899 (2405) GTTCGTAGTAAGAGAAATCTTCTGCCTTGAGATGCTGTTAA-----

ENV GENOMIC HERV-KII (1850) GTTCGTAGTAA-----

ENV GENOMIC AC008813 (1238) -----

ENV GENOMIC AC012309 (2133) GTTCGTAGCAAG-----

ENV GENOMIC AL121932 (1538) -----

ENV GENOMIC AD000090 (2157) GTTCGTAGTAA-----

ENV GEN AL160008 (647) -----

ENV GENOMIC HEU32496 (441) -----

ENV GENOMIC AC011467 (1699) -----

ENV GENOMIC AF235103 (2212) GAAAAAGAGCTGTACTTTGAACAATTGCTTGTCTCAGATGTTGTTAATTTGTAGTTTT-----

ENV GENOMIC AC026786 (2086) GAAAAAGAGCTGTACTTTAAACAATTGCTTGTCTCAGATGTTGTTAATTTGTAGCTTTCCCGAGCC-----

ENV GENOMIC AC034203 (1403) -----

ENV GENOMIC AC018809 (1846) GTCTCGTGTAGTAAG-----

ENV GENOMIC HERV-KI02 AF164610 (1906) GTTTAGTGTGTAAGAAAAATCTT-----

ENV GENOMIC FRAG. AF260253 (385) GTTCGTAGTAA-----

CONSENSUS (2641) TG TGTAC

FIGURE 7

1 60

GI_4185938_EMB_CAA76878.1_ (1) -----MGQTKSKIISKYASYLSFIKILLKRGVVKVSTKNLIKLFQIIIEQFCPWFPEQGT

GI_4185942_EMB_CAA76881.1_ (1) -----MGQTKSKIISKYASYLSFIKILLKRGVVKVSTKNLIKLFQIIIEQFCPWFPEQGT

GI_4185946_EMB_CAA76884.1_ (1) -----MGQTKSKIISKYASYLSFIKILLKRGVVKVSTKNLIKLFQIIIEQFCPWFPEQGT

GI_5931704_EMB_CAB56602.1_ (1) -----MGQTKTKSKYASYLSFIKILLKRGVVRVSTKNLIKLFQIIIEQFCPWFPEQGT

GAG OF AB047240 (1) -----MGQTKSKTKSKYASYLSFIKILLKRGVVRVSTKNLIKLFQIIIEQFCPWFPEQGT

TRANSLATION OF ORF99 (1) YKAGLGQTKSKTKSKYASYLSFIKILLKRGVVRVSTKNLIKLFQIIIEQFCPWFPEQGT

TRANSLATION OF G226TOP-LINK (1) -----

TRANSLATION OF G591TOP-LINK (1) -----

TRANSLATION OF LNCAP-GAG (1) -----MGQTKSKTKSKYASYLSFIKILLKRGVVRVSTKNLIKLFQIIIEQFCPWFPEQGT

GAG106-135 (1) -----

GAG186-215 (1) -----

GAG46-75 (1) -----CPWFPEQGT

PDG-G1 (1) -----

PGD-G2 (1) -----

PGD-G3 (1) -----

CONSENSUS (1) CPWFPEQG L

61 120

GI_4185938_EMB_CAA76878.1_ (56) DLKDWKRIGKELKQAGRKGNIPLTVWNDWAIKAALPEPQTEDSVSVSDAPGSCI IDC

GI_4185942_EMB_CAA76881.1_ (56) DLKDWKRIGKELKQAGRKGNIPLTVWNDWAIKAALPEPQTEDSVSVSDAPGSCI IDC

GI_4185946_EMB_CAA76884.1_ (56) DLKDWKRIGKELKQAGRKGNIPLTVWNDWAIKAALPEPQTEDSVSVSDAPGSCI IDC

GI_5931704_EMB_CAB56602.1_ (54) DLEDWKRIGKELKQAGRKGNIPLTVWNDWPIKAALPEPQTEDS-VSVSDAPGSCI IDC

GAG OF AB047240 (56) DLKDWKRIGBELKQAGRKGNIPLTVWNDWAIKAALPEPQTEDSVSVSDAPGSCI IDC

TRANSLATION OF ORF99 (61) DLKDWKRIGBELKQAGRKGNIPLTVWNDWAIKAALPEPQTEDSVSVSDAPGSCI IDC

TRANSLATION OF G226TOP-LINK (1) -----

TRANSLATION OF G591TOP-LINK (1) -----

TRANSLATION OF LNCAP-GAG (56) DLKDWKRIGBELKQAGRKGNIPLTVWNDWAIKAALPEPQTEDSVSVSDAPGSCI IDC

GAG106-135 (1) -----DAPGSCI IDC

GAG186-215 (1) -----

GAG46-75 (11) DLKDWKRIGKELKQAGRKGNI-----

PDG-G1 (1) ---DWKRIGKELKQAGRKG-----

PGD-G2 (1) -----

PGD-G3 (1) -----

CONSENSUS (61) DL DWKRIG ELKQAGRKGNI DAPGSCI IDC

FIGURE 7 CONTD...

GI_4185938_EMB_CAA76878.1_	(116)	NENTR	KKSQKETEGHCEYVAEPVMAQSTQNVNDYNQLQEVIYPETLKLKLGKGPPELVGPSE	121	180
GI_4185942_EMB_CAA76881.1_	(116)	NENTR	KKSQKETESLHCEYVAEPVMAQSTQNVNDYNQLQEVIYPETLKLKLGKGPPELVGPSE		
GI_4185946_EMB_CAA76884.1_	(116)	NENTR	KKSQKETEGHCEYVAEPVMAQSTQNVNDYNQLQEVIYPETLKLKLGKGPPELVGPSE		
GI_5931704_EMB_CAB56602.1_	(113)	NEKTR	KKSQKETETLHCEYVAEPLMAQSTQNVNDYNQLQEVIYPETLKLKLGKGPPELVGPSE		
GAG OF AB047240	(116)	NEKTGR	KKSQKETESLHCEYVTEPVMAQSTQNVNDYNQLQGVYIPETLKLKLGKGPPELVGPSE		
TRANSLATION OF ORF99	(121)	NEKTGR	KKSQKETESLHCEYVTEPVMAQSTQNVNDYNQLQGVYIPETLKLKLGKGPPELVGPSE		
TRANSLATION OF G226TOP-LINK	(1)		-----		
TRANSLATION OF G591TOP-LINK	(1)		-----		
TRANSLATION OF LNCAP-GAG	(116)	NEKTGR	KKSQKETESLHCEYVTEPVMAQSTQNVNDYNQLQGVYIPETLKLKLGKGPPELVGPSE		
GAG106-135	(11)	NENTR	KKSQKETEGHCEYV-----		
GAG186-215	(1)		-----		
GAG46-75	(31)		-----		
PDG-G1	(17)		-----		
PGD-G2	(1)		-----		
PGD-G3	(1)		-----		
CONSENSUS	(121)	NE T	KKSQKETE LHCEYV		
GI_4185938_EMB_CAA76878.1_	(176)	SKPRGTSPLPAGQV	PVTLQPQKQV	181	240
GI_4185942_EMB_CAA76881.1_	(176)	SKPRGTSRLPAGQV	PVTLQPQKQV		
GI_4185946_EMB_CAA76884.1_	(176)	SKPRGTSPLPAGQV	PVTLQPQKQV		
GI_5931704_EMB_CAB56602.1_	(173)	SKPRGPSPLSAGQV	TVTLQPQKQV		
GAG OF AB047240	(176)	SKPRGPSPLPAGQV	PVTLQPQKQV		
TRANSLATION OF ORF99	(181)	SKPRGPSPLPAGQV	PVTLQPQKQV		
TRANSLATION OF G226TOP-LINK	(1)		-----		
TRANSLATION OF G591TOP-LINK	(1)		-----		
TRANSLATION OF LNCAP-GAG	(176)	SKPRGPSPLPAGQV	PVTLQPQKQV		
GAG106-135	(31)		-----		
GAG186-215	(1)		-----		
GAG46-75	(31)		-----		
PDG-G1	(17)		-----		
PGD-G2	(1)		-----		
PGD-G3	(1)		-----		
CONSENSUS	(181)	AGQV	VTLPQK QV	KENKTQ	PVAYQYWPP SQYGY GMPP
GI_4185938_EMB_CAA76878.1_	(236)	APQGRAPYPQPPT	TRLNPTAPP	241	300
GI_4185942_EMB_CAA76881.1_	(236)	APQGRAPYPQPPT	TRLNPTAPP		
GI_4185946_EMB_CAA76884.1_	(236)	APQGRAPYPQPPT	TRLNPTAPP		
GI_5931704_EMB_CAB56602.1_	(233)	APQDREPYPQPPT	TRRQCYGTT		
GAG OF AB047240	(236)	ALQGRAPYPQPPT	TRLNPTASRSGQGGT		
TRANSLATION OF ORF99	(241)	ALQGRAPYPQPPT	TRLNPTASRSGQGGT		
TRANSLATION OF G226TOP-LINK	(11)	APQGRAPYPQPPT	TRLNPTA		
TRANSLATION OF G591TOP-LINK	(1)		-----		
TRANSLATION OF LNCAP-GAG	(236)	ALQGRAPYPQPPT	TRLNPTASRSGQGGT		
GAG106-135	(31)		-----		
GAG186-215	(31)		-----		
GAG46-75	(31)		-----		
PDG-G1	(17)		-----		
PGD-G2	(1)		-----		
PGD-G3	(1)		-----		
CONSENSUS	(241)	A Q R	PYPQPPT R		
GI_4185938_EMB_CAA76878.1_	(296)	QEGEPPTVEARYK	SFSIKKLDKMEGVKQYGN	301	360
GI_4185942_EMB_CAA76881.1_	(296)	QEGEPPTVEARYK	SFSIKKLDKMEGVKQYGN		
GI_4185946_EMB_CAA76884.1_	(296)	QEGEPPTVEARYK	SFSIKKLDKMEGVKQYGN		
GI_5931704_EMB_CAB56602.1_	(254)		-----		
GAG OF AB047240	(296)	QVGAPARAETRC	EPFTMKMLDKIKEGVKQYGSNSPY		
TRANSLATION OF ORF99	(301)	QVGAPARAETRC	EPFTMKMLDKIKEGVKQYGSNSPY		
TRANSLATION OF G226TOP-LINK	(31)		-----		
TRANSLATION OF G591TOP-LINK	(1)		-----		
TRANSLATION OF LNCAP-GAG	(296)	QVGAPARAETRC	EPFTMKMLDKIKEGVKQYGSNSPY		
GAG106-135	(31)		-----		
GAG186-215	(31)		-----		
GAG46-75	(31)		-----		
PDG-G1	(17)		-----		
PGD-G2	(17)		-----		
PGD-G3	(1)		-----		
CONSENSUS	(301)		-----		

	361	420
GI_4185938_EMB_CAA76878.1_	(356)	SSLSPSQFLQFKTWWIDGVQEQVRRNRANPPVNIADADQLLGIGQNWSTISQQALMQNEA
GI_4185942_EMB_CAA76881.1_	(356)	SSLSPSQFLQFKTWWIDGVQEQVRRNRANPPVNIADADQLLGIGQNWSTISQQALMQNEA
GI_4185946_EMB_CAA76884.1_	(356)	SSLSPSQFLQFKTWWIDGVQEQVRRNRANPPVNIADADQLLGIGQNWSTISQQALMQNEA
GI_5931704_EMB_CAB56602.1_	(254)	
GAG OF AB047240	(356)	SSLSSSQYLQFKTWWIDGVQEQVRKNQATKPTVNIADADQLLGTGPNWSTINQQSVMQNEA
TRANSLATION OF ORF99	(361)	SSLSSSQYLQFKTWWIDGVQEQVRKNQATKPTVNIADADQLLGTGPNWSTINQQSVMQNEA
TRANSLATION OF G226TOP-LINK	(31)	-----
TRANSLATION OF G591TOP-LINK	(1)	-----
TRANSLATION OF LNCAP-GAG	(356)	SSLSSSQYLQFKTWWIDGVQEQVRKNQATKPTVNIADADQLLGTGPNWSTINQQSVMQNEA
GAG106-135	(31)	-----
GAG186-215	(31)	-----
GAG46-75	(31)	-----
PDG-G1	(17)	-----
PGD-G2	(17)	-----
PGD-G3	(1)	-----
CONSENSUS	(361)	
	421	480
GI_4185938_EMB_CAA76878.1_	(416)	IEQVRAICLRAWEKIQDPGSTCPSFNTVRQGSKEPYPDFVARLQDVAQKSIADKARKVI
GI_4185942_EMB_CAA76881.1_	(416)	IEQVRAICLRAWEKIQDPGSTCPSFNTVRQGSKEPYPDFVARLQDVAQKSIANERKARKVI
GI_4185946_EMB_CAA76884.1_	(416)	IEQVRAICLRAWEKIQDPGSTCPSFNTVRQGSKEPYPDFVARLQDVAQKSIADKARKVI
GI_5931704_EMB_CAB56602.1_	(254)	
GAG OF AB047240	(416)	IEQVRAICLRAWGKIQDPGTAF - INSIRQGSKEPYPDFVARLQDAAQKSITDDNARKVI
TRANSLATION OF ORF99	(421)	IEQVRAICLRAWGKIQDPGTAF - PINSIRQGSKEPYPDFVARLQDAAQKSITDDNARKVI
TRANSLATION OF G226TOP-LINK	(31)	-----
TRANSLATION OF G591TOP-LINK	(1)	-----
TRANSLATION OF LNCAP-GAG	(416)	IEQVRAICLRAWGKIQDPGTAF - INSIRQGSKEPYPDFVARLQDAAQKSITDDNARKVI
GAG106-135	(31)	-----
GAG186-215	(31)	-----
GAG46-75	(31)	-----
PDG-G1	(17)	-----
PGD-G2	(17)	-----
PGD-G3	(1)	-----
CONSENSUS	(421)	
	481	540
GI_4185938_EMB_CAA76878.1_	(476)	VELMAYENANPECQSAIKPLKGKVPAGSDVISEYVKACDGIGGAMYKAMLMAQAITGVVL
GI_4185942_EMB_CAA76881.1_	(476)	VELMAYENPNPECQSAIKPLKGKVPAGSDVISEYVKACDGMGGAMHKAMLMAQAITGVVL
GI_4185946_EMB_CAA76884.1_	(476)	VELMAYENANPECQSAIKPLKGKVPAGSDVISEYVKACDGIGGAMHKAMLMAQAITGVVL
GI_5931704_EMB_CAB56602.1_	(254)	
GAG OF AB047240	(475)	VELMAYENANPECQSAIKPLKGKVPAGVDVITEYVKACDGIGGAMHKAMLMAQAMRGLTL
TRANSLATION OF ORF99	(480)	VELMAYENANPECQSAIKPLKGKVPAGVDVITEYVKACDGIGGAMHKAMLMAQAMRGLTL
TRANSLATION OF G226TOP-LINK	(31)	-----
TRANSLATION OF G591TOP-LINK	(1)	-----
TRANSLATION OF LNCAP-GAG	(475)	VELMAYENANPECQSAIKPLKGKVPAGVDVITEYVKACDGIGGAMHKAMLMAQAMRGLTL
GAG106-135	(31)	-----
GAG186-215	(31)	-----
GAG46-75	(31)	-----
PDG-G1	(17)	-----
PGD-G2	(17)	-----
PGD-G3	(1)	-----
CONSENSUS	(481)	
	541	600
GI_4185938_EMB_CAA76878.1_	(536)	GGQVRTFGKRCYNCGQIGHLKKNCPLVNKQNIITIQATTG - REPPDLCPRCCKGKHWASQ
GI_4185942_EMB_CAA76881.1_	(536)	GGQVRTFGGKRCYNCGQIGHLKKNCPLVNKQNIITIQATTG - REPPDLCPRCCKGKHWASQ
GI_4185946_EMB_CAA76884.1_	(536)	GGQVRTFGKRCYNCGQIGHLKKNCPLVNKQNIITIQATTG - REPPDLCPRCCKGKHWASQ
GI_5931704_EMB_CAB56602.1_	(254)	
GAG OF AB047240	(535)	GGQVRTFGKKCYNCGQIGHLKRSCPLVNKQNIINQAITAKNKKPSGLCPKCGKGKHWANQ
TRANSLATION OF ORF99	(540)	GGQVRTFGKKCYNCGQIGHLKRSCPLVNKQNIINQAITAKNKKPSGLCPKCGKGKHWANQ
TRANSLATION OF G226TOP-LINK	(31)	-----
TRANSLATION OF G591TOP-LINK	(1)	-----
TRANSLATION OF LNCAP-GAG	(535)	GGQVRTFGKKCYNCGQIGHLKRSCPLVNKQNIINQAITAKNKKPSGLCPKCGKGKHWANQ
GAG106-135	(31)	-----
GAG186-215	(31)	-----
GAG46-75	(31)	-----
PDG-G1	(17)	-----
PGD-G2	(17)	-----
PGD-G3	(1)	-----
CONSENSUS	(541)	

FIGURE 7 CONTD...

	601	660
GI_4185938_EMB_CAA76878.1_	(595)	CRSKFDKNGQPLSGNEQRGQPAPQQTGAFFIQPFVPGFQGGQP-PLSQVFQGISQLPQ
GI_4185942_EMB_CAA76881.1_	(595)	CRSKFDKNGQPLSGNEQRGQPAPQQTGAFFIQPFVPHGFQGGQP-PLSQVFQGISQLPQ
GI_4185946_EMB_CAA76884.1_	(595)	CRSKFDKNGQPLSGNEQRGQPAPQQTGAFFIQPFVPGFQGGQP-PLSQVFQGISQLPQ
GI_5931704_EMB_CAB56602.1_	(254)	-----
GAG OF AB047240	(595)	CHSKFDKDGQPLSGNRKRGPQAPQQTGAFFVQLFVPQGFQGGQPLQKIPPLQGVSQLQQ
TRANSLATION OF ORF99	(600)	CHSKFDKDGQPLSGNRKRGPQAPQQTGAFFVQLFVPQGFQGGQPLQKIPPLQGVSQLQQ
TRANSLATION OF G226TOP-LINK	(31)	-----
TRANSLATION OF G591TOP-LINK	(5)	CRSKFDKNGQPLSGNEQRGQPAPQ-----
TRANSLATION OF LNCAP-GAG	(595)	CHSKFDKDGQPLSGNRKRGPQAPQQTGAFFVQLFVPQGFQGGQPLQKIPPLQGVSQLQQ
GAG106-135	(31)	-----
GAG186-215	(31)	-----
GAG46-75	(31)	-----
PDG-G1	(17)	-----
PGD-G2	(17)	-----
PGD-G3	(1)	CRSKFDKNGQPLSGNE-----
CONSENSUS	(601)	C SKFDK GQPLSGN

	661	673
GI_4185938_EMB_CAA76878.1_	(654)	YNNCPPPQAAVQQ
GI_4185942_EMB_CAA76881.1_	(654)	YNNCPPPQAAVQQ
GI_4185946_EMB_CAA76884.1_	(654)	YNNCPPPQAAVQQ
GI_5931704_EMB_CAB56602.1_	(254)	-----
GAG OF AB047240	(655)	SNSCPAPQQAAPQ
TRANSLATION OF ORF99	(660)	SNSCPAPQQAAPQ
TRANSLATION OF G226TOP-LINK	(31)	-----
TRANSLATION OF G591TOP-LINK	(31)	-----
TRANSLATION OF LNCAP-GAG	(655)	SNSCPAPQQAAPQ
GAG106-135	(31)	-----
GAG186-215	(31)	-----
GAG46-75	(31)	-----
PDG-G1	(17)	-----
PGD-G2	(17)	-----
PGD-G3	(17)	-----
CONSENSUS	(661)	

FIGURE 8

	1	60
GI_4185939_EMB_CAA76879.1_	(1)	MLTDLRAVN---AVIQPMGPLQPLGSPAMIPKDWPLIIIDLKDCFFTIPLAEQDCEKFA
GI_4185943_EMB_CAA76882.1_	(1)	MLTDLRAVNAVNAVIVQPMGPLQPLGSLAMIPKDWPLIIIDLKDCFFTIPLAEQDCEKFA
GI_4185947_EMB_CAA76885.1_	(1)	MLTDLRAVN---AVIQPMGPLQPLGSPAMIPKDWPLIIIDLKDCFFTIPLAEQDCEKFA
GI_5931705_EMB_CAB56603.1_	(1)	-----MIPKDWPLIIIDLKDCFFTIPLAEQDCEKFA
ENV OF AB047240	(1)	-----
TRANSLATION OF P386TOP-LINK	(1)	-----
TRANSLATION OF POL349-LINK	(1)	-----
LNCAP-GENOMEA-POLORF	(1)	-----
TRANSLATION OF LNCAP-POL-GENA-GOODA	(1)	-----
TRANSLATION OF ORF111-10	(1)	-----
PGD-P1	(1)	-----
PGD-P2	(1)	-----
PGDP3	(1)	-----
CONSENSUS	(1)	

	61	120
GI_4185939_EMB_CAA76879.1_	(58)	FTIPAINNKEPATRFQWKVLPQGMNSPTICQTFVGRALQPVREKFSDCYIIHCIDDILC
GI_4185943_EMB_CAA76882.1_	(61)	FTIPAINNKEPATRFQWKVLPQGMNSPTICQTFVGRALQPVREKFSDCYIIHYIDDILC
GI_4185947_EMB_CAA76885.1_	(58)	FTIPAINNKEPATRFQWKVLPQGMNSPTICQTFVGRALQPVREKFSDCYIIHCIDDILC
GI_5931705_EMB_CAB56603.1_	(32)	FTIPAINNKEPATRFQWKVLPQGMNSPTLCQTFVGRALQPVREKFSDCYIIHYFDDILC
ENV OF AB047240	(1)	-----
TRANSLATION OF P386TOP-LINK	(1)	-----
TRANSLATION OF POL349-LINK	(1)	-----
LNCAP-GENOMEA-POLORF	(1)	-----
TRANSLATION OF LNCAP-POL-GENA-GOODA	(1)	-----
TRANSLATION OF ORF111-10	(1)	-----
PGD-P1	(1)	-----
PGD-P2	(1)	-----
PGDP3	(1)	-----
CONSENSUS	(61)	

121 140

GI_4185939_EMB_CAA76879.1 (118) A A E T K D K L I D C Y T F L Q A E V A N A G L A I A S D K I Q T S T P P H Y L G M Q I E N R K I K P Q K I E I R K D T
 GI_4185943_EMB_CAA76882.1 (121) A A E M K D K L I D C Y T F L Q A E V A N A G L A I A S D K I Q T S T P P H Y L G M Q I E N R K I K P Q K I E I R K D T
 GI_4185947_EMB_CAA76885.1 (118) A A E T K D K L I D C Y T F L Q A E V A N A G L A I A S D K I Q T S T P P H Y L G M Q I E N R K I K P Q K I E I R K D T
 GI_5931705_EMB_CAB56603.1 (92) A A E T K D K L I D C Y T F L Q A E V A N A G L A I A S D K I Q T S T P P H Y L G M Q I E N R K I K P Q K I E I R K D T
 ENV OF AB047240 (1) -----
 TRANSLATION OF P386TOP-LINK (1) -----
 TRANSLATION OF POL349-LINK (1) -----
 LNCAP-GENOMEA-POLORF (1) -----
 TRANSLATION OF LNCAP-POL-GENA-GOODA (1) -----
 TRANSLATION OF ORF111-10 (1) -----
 PGD-P1 (1) -----
 PGD-P2 (1) -----
 PGDP3 (1) -----
 CONSENSUS (121) -----

181 240

GI_4185939_EMB_CAA76879.1 (178) L K T L N D F Q K L L G D I N W I R P T L G I P T Y A M S N L F S I L R G D S D L N S K R M L T P E A T K E I K L V E E
 GI_4185943_EMB_CAA76882.1 (181) L K T L N D F Q K L L G D I N W I R P T L G I P T Y A M S N L F S I L R G D S D L N S K R M L T P E A T K E I K L V E E
 GI_4185947_EMB_CAA76885.1 (178) L K T L N D F Q K L L G D I N W I R P T L G I P T Y A M S N L F S I L R G D S D L N S K R M L T P E A T K E I K L V E E
 GI_5931705_EMB_CAB56603.1 (152) L K T L N D F Q K L L G D I N W I R P T L G I P T Y A M S N L F S I L R G D S D L N S K R M L T P E A T K E I K L V E E
 ENV OF AB047240 (1) -----
 TRANSLATION OF P386TOP-LINK (1) -----
 TRANSLATION OF POL349-LINK (1) -----
 LNCAP-GENOMEA-POLORF (1) -----
 TRANSLATION OF LNCAP-POL-GENA-GOODA (1) -----
 TRANSLATION OF ORF111-10 (1) -----
 PGD-P1 (17) -----
 PGD-P2 (1) -----
 PGDP3 (1) -----
 CONSENSUS (181) -----

241 300

GI_4185939_EMB_CAA76879.1 (238) K I Q S A Q I N R I D P L A P L Q L I F T A H S P T I I I Q N T D L V W S F L P H S T K T F T L Y L D Q M A T
 GI_4185943_EMB_CAA76882.1 (241) K I Q S A Q I N R I D P L A P L Q L I F T A H S P T I I I Q N T D L V W S F L P H S T K T F T L Y L D Q M A T
 GI_4185947_EMB_CAA76885.1 (238) K I Q S A Q I N R I D P L A P L Q L I F T A H S P T I I I Q N T D L V W S F L P H S T K T F T L Y L D Q M A T
 GI_5931705_EMB_CAB56603.1 (212) K I Q S A Q I N R I D P L A P L Q L I F T A H S P T I I I Q N T D L V W S F L P H S T K T F T L Y L D Q M A T
 ENV OF AB047240 (1) -----
 TRANSLATION OF P386TOP-LINK (1) -----
 TRANSLATION OF POL349-LINK (1) -----
 LNCAP-GENOMEA-POLORF (1) -----
 TRANSLATION OF LNCAP-POL-GENA-GOODA (1) -----
 TRANSLATION OF ORF111-10 (1) -----
 PGD-P1 (17) -----
 PGD-P2 (1) -----
 PGDP3 (1) -----
 CONSENSUS (241) D L A P L Q L L I F A T A H S T G I I I Q N T D L V W S F L P H S T K T F T L Y L D Q M A T

301 360

GI_4185939_EMB_CAA76879.1 (298) L I G Q T R L R I I K L C G N D P D K I V P F N K Q V R Q A F I N S G A W Q I G L A N F I G I I D N H Y P K T K I F
 GI_4185943_EMB_CAA76882.1 (301) L I G Q T R L R I I K L C G N D P D K I V P F N K Q V R Q A F I N S G A W Q I G L A N F I G I I D N H Y P K T K I F
 GI_4185947_EMB_CAA76885.1 (298) L I G Q T R L R I I K L C G N D P D K I V P F N K Q V R Q A F I N S G A W Q I G L A N F I G I I D N H Y P K T K I F
 GI_5931705_EMB_CAB56603.1 (272) L I G P T R L R I I K L C G N D P D K I V P F N K Q V R Q A F I N S G A W Q I G L A N F I G I I D N H Y P K T K I F
 ENV OF AB047240 (4) L I G Q G R L R I I T L C G N D P D K I T V P F N K Q V R Q A F I S S G A W Q I G L A N F I G I I D N H Y P K T K I F
 TRANSLATION OF P386TOP-LINK (1) -----
 TRANSLATION OF POL349-LINK (1) -----
 LNCAP-GENOMEA-POLORF (1) -----
 TRANSLATION OF LNCAP-POL-GENA-GOODA (51) L I G Q G R L R I I T L C G N D P D K I T V P F N K Q V R Q A F I S S G A W Q I G L A N F I G I I D N H Y P K T K I F
 TRANSLATION OF ORF111-10 (57) L I G Q G R L R I I T L C G N D P D K I T V P F N K Q V R Q A F I S S G A W Q I G L A N F I G I I D N H Y P K T K I F
 PGD-P1 (17) -----
 PGD-P2 (1) -----
 PGDP3 (1) -----
 CONSENSUS (301) L I G Q R L R I I L C G N D P D K I V P F N K Q V R Q A F I S S G A W Q I G L A N F I G I I D N H Y P K T K I F

361 420

GI_4185939_EMB_CAA76879.1 (358) Q F L K L T T W I L P K I T R R E P L E N A L T V F T D G S S N G K A A Y T G P K E R V I K T P Y O S A Q R A E L V A V
 GI_4185943_EMB_CAA76882.1 (361) Q F L K L T T W I L P K I T R R E P L E N A L T V F T D G S S N G K A A Y T G P K E R V I K T P Y O S A Q R A E L V A V
 GI_4185947_EMB_CAA76885.1 (358) Q F L K L T T W I L P K I T R R E P L E N A L T V F T D G S S N G K A A Y T G P K E R V I K T P Y O S A Q R A E L V A V
 GI_5931705_EMB_CAB56603.1 (332) Q F L K L T T W I L P K I T R R E P L E N A L T V F T D G S S N G K A A Y T G P K E R V I K T P Y O S A Q R A E L V A V
 ENV OF AB047240 (64) Q F L K L T T W I L P K I T R R E P L E N A L T V F T D G S S N G K A A Y T G P K E R V I K T P Y O S A Q R A E L V A V
 TRANSLATION OF P386TOP-LINK (1) -----
 TRANSLATION OF POL349-LINK (10) Q F L K L T T W I L P K I T R R E P - - - - - G S S N G K A A Y T G P K E R V I K T P Y O S A Q R A E L V - -
 LNCAP-GENOMEA-POLORF (111) Q F L K L T T W I L P K I T R R E P L E N A L T V F T D G S S N G K A A Y T G P K E R V I K T P Y O S A Q R A E L V A V
 TRANSLATION OF LNCAP-POL-GENA-GOODA (111) Q F L K L T T W I L P K I T R R E P L E N A L T V F T D G S S N G K A A Y T G P K E R V I K T P Y O S A Q R A E L V A V
 TRANSLATION OF ORF111-10 (117) Q F L K L T T W I L P K I T R R E P L E N A L T V F T D G S S N G K A A Y T G P K E R V I K T P Y O S A Q R A E L V A V
 PGD-P1 (17) -----
 PGD-P2 (1) -----
 PGDP3 (1) -----
 CONSENSUS (361) Q F L K L T T W I L P K I T R R E P L E N A L T V F T D G S S N G K A A Y T G P K E R V I K T P Y O S A Q R A E L V A V

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FIGURE 8 contd...

GI_4185939_EMB_CAA76879.1	(418)	ITVLQDFDQPINIISDSAYVVQATRDVETALIKYSMDQDLNQLFNLLQOTVRKRNFPPFYI	421	480
GI_4185943_EMB_CAA76882.1	(421)	ITVLQDFDQPINIISDSAYVVQATRDVETALIKYSMDQDLNQLFNLLQOTVRKRNFPPFYI		
GI_4185947_EMB_CAA76885.1	(418)	ITVLQDFDQPINIISDSAYVVQATRDVETALIKYSMDQDLNQLFNLLQOTVRKRNFPPFYI		
GI_5931705_EMB_CAB56603.1	(392)	ITVLQDFDQPINIISDSAYVVQATRDVETALIKYSMDQDLNQLFNLLQOTVRKRNFPPFYI		
ENV OF AB047240	(124)	ITVLQDFDQPINIISDSAYVVQATRDVETALIKYSTDDHLNQLFNLLQOTVRKRNFPPFYI		
TRANSLATION OF P386TOP-LINK	(31)	-----		
TRANSLATION OF POL349-LINK	(28)	-----		
LNCAP-GENOMEA-POLORF	(171)	ITVLQDFDQPINIISDSAYVVQATRDVETALIKYSTDDHLNQLFNLLQOTVRKRNFPPFYI		
TRANSLATION OF LNCAP-POL-GENA-GOODA	(171)	ITVLQDFDQPINIISDSAYVVQATRDVETALIKYSTDDHLNQLFNLLQOTVRKRNFPPFYI		
TRANSLATION OF ORF111-10	(177)	ITVLQDFDQPINIISDSAYVVQATRDVETALIKYSTDDHLNQLFNLLQOTVRKRNFPPFYI		
PGD-P1	(17)	-----		
PGD-P2	(17)	-----		
PGDP3	(1)	-----		
CONSENSUS	(421)	ITVLQDFDQPINIISDSAYVVQATRDVETALIKYS DD LNQLFNLLQOTVRKRNFPPFYI		
GI_4185939_EMB_CAA76879.1	(478)	THIRAHTNLPGLPTKANEQADLLVSSALIKAEQLHALTHVNAAGLNKPKFVDTWKQAKDIV	481	540
GI_4185943_EMB_CAA76882.1	(481)	THIRAHTNLPGLPTKANEQADLLVSSALIKAEQLHALTHVNAAGLNKPKFVDTWKQAKDIV		
GI_4185947_EMB_CAA76885.1	(478)	THIRAHTNLPGLPTKANEQADLLVSSALIKAEQLHALTHVNAAGLNKPKFVDTWKQAKDIV		
GI_5931705_EMB_CAB56603.1	(452)	THIRAHTNLPGLPTKANEQADLLVSSAFIKAEQLHALTHVNAAGLNKPKFVDTWKQAKDIV		
ENV OF AB047240	(184)	THIRAHTNLPGLPTKANEQADLLVSSAFIKAEQLHALTHVNAAGLNKPKFVDTWKQAKDIV		
TRANSLATION OF P386TOP-LINK	(31)	-----		
TRANSLATION OF POL349-LINK	(28)	-----		
LNCAP-GENOMEA-POLORF	(231)	THIRAHTNLPGLPTKANEQADLLVSSAFIKAEQLHALTHVNAAGLNKPKFVDTWKQAKDIV		
TRANSLATION OF LNCAP-POL-GENA-GOODA	(231)	THIRAHTNLPGLPTKANEQADLLVSSAFIKAEQLHALTHVNAAGLNKPKFVDTWKQAKDIV		
TRANSLATION OF ORF111-10	(237)	THIRAHTNLPGLPTKANEQADLLVSSAFIKAEQLHALTHVNAAGLNKPKFVDTWKQAKDIV		
PGD-P1	(17)	-----		
PGD-P2	(17)	-----		
PGDP3	(1)	-----		
CONSENSUS	(481)	THIRAHTNLPGLPTKANEQADLLVSSA IKAQEL ALTHVNAAGLNKPKFVDTWKQAKDIV		
GI_4185939_EMB_CAA76879.1	(538)	QHCTQCQVHLHLPQAEAGVNPRLGCPNALWQMDVTHVPSFGRLSYVHVTVDTYSHFIWATC	541	600
GI_4185943_EMB_CAA76882.1	(541)	QHCTQCQVHLHLPQAEAGVNPRLGCPNALWQMDVTHVPSFGRLSYVHVTVDTYSHFIWATC		
GI_4185947_EMB_CAA76885.1	(538)	QHCTQCQVHLHLPQAEAGVNPRLGCPNALWQMDVTHVPSFGRLSYVHVTVDTYSHFIWATC		
GI_5931705_EMB_CAB56603.1	(512)	QHCTQCQVLDLPQAEAGVNPVCLMHYGKWMSHMYLHLGRLSYVHVTVDTYSHFIWATC		
ENV OF AB047240	(244)	QHCTQCQVHLHLPQAEAGVNPRLGCPNALWQMDGTHVPSFGRLSYVHVTVDTYSHFIWATC		
TRANSLATION OF P386TOP-LINK	(31)	-----		
TRANSLATION OF POL349-LINK	(28)	-----		
LNCAP-GENOMEA-POLORF	(291)	QHCTQCQVHLHLPQAEAGVNPRLGCPNALWQMDGTHVPSFGRLSYVHVTVDTYSHFIWATC		
TRANSLATION OF LNCAP-POL-GENA-GOODA	(291)	QHCTQCQVHLHLPQAEAGVNPRLGCPNALWQMDGTHVPSFGRLSYVHVTVDTYSHFIWATC		
TRANSLATION OF ORF111-10	(297)	QHCTQCQVHLHLPQAEAGVNPRLGCPNALWQMDGTHVPSFGRLSYVHVTVDTYSHFIWATC		
PGD-P1	(17)	-----		
PGD-P2	(17)	-----		
PGDP3	(1)	-----		
CONSENSUS	(541)	QHCTQCQVHLHLPQAEAGVNPRLGCPNALWQMD THV SFGRLSYVHVTVDTYSHFIWATC		
GI_4185939_EMB_CAA76879.1	(598)	QTGESTSHVKKHLLSCFAVMGVPEKIKTDNGPGYCSKAFQKFLSOWKISHTTGIPYNSQG	601	660
GI_4185943_EMB_CAA76882.1	(601)	QTGESTSHVKKHLLSCFAVMGVPEKIKTDNGPGYCSKAFQKFLSOWKISHTTGIPYNSQG		
GI_4185947_EMB_CAA76885.1	(598)	QTGESTSHVKKHLLSCFAVMGVPEKIKTDNGPGYCSKAFQKFLSOWKISHTTGIPYNSQG		
GI_5931705_EMB_CAB56603.1	(572)	QTGESTSHVKKHLLSCFAVMGVPEKIKTDNGPGYCSKAFQKFLSOWKISHTTGIPYNSQG		
ENV OF AB047240	(304)	QTGESTSHVKKHLLSCFAVMGVPEKIKTDNGPGYCSKAFQKFLSOWKISHTTGIPYNSQG		
TRANSLATION OF P386TOP-LINK	(31)	-----		
TRANSLATION OF POL349-LINK	(28)	-----		
LNCAP-GENOMEA-POLORF	(351)	QTGESTSHVKKHLLSCFAVMGVPEKIKTDNGPGYCSKAFQKFLSOWKISHTTGIPYNSQG		
TRANSLATION OF LNCAP-POL-GENA-GOODA	(351)	QTGESTSHVKKHLLSCFAVMGVPEKIKTDNGPGYCSKAFQKFLSOWKISHTTGIPYNSQG		
TRANSLATION OF ORF111-10	(357)	QTGESTSHVKKHLLSCFAVMGVPEKIKTDNGPGYCSKAFQKFLSOWKISHTTGIPYNSQG		
PGD-P1	(17)	-----		
PGD-P2	(17)	-----		
PGDP3	(1)	-----		
CONSENSUS	(601)	QTGESTSHVKKHLLSCFAVMGVPEKIKTDNGPGYCSKAFQKFLSOWKISHTTGIPYNSQG		
GI_4185939_EMB_CAA76879.1	(658)	QAIVERTNRTLKTQLVKQKEGGDSKECTTPQMQLNLALYTLNPLNIYRNQTTTSAQOHLT	661	720
GI_4185943_EMB_CAA76882.1	(661)	QAIVERTNRTLKTQLVKQKEGGDSKECTTPQMQLNLALYTLNPLNIYRNQTTTSAQOHLT		
GI_4185947_EMB_CAA76885.1	(658)	QAIVERTNRTLKTQLVKQKEGGDSKECTTPQMQLNLALYTLNPLNIYRNQTTTSAQOHLT		
GI_5931705_EMB_CAB56603.1	(632)	QAIVERTNRTLKTQLVKQKEGGDSKECTTPQMQLNLALYTLNPLNIYRNQTTTSAQOHLT		
ENV OF AB047240	(364)	QAIVERTNRTLKTQLVKQKEGGDSKECTTPQMQLNLALYTLNPLNIYRNQTTTSAQOHLT		
TRANSLATION OF P386TOP-LINK	(31)	-----		
TRANSLATION OF POL349-LINK	(28)	-----		
LNCAP-GENOMEA-POLORF	(411)	QAIVERTNRTLKTQLVKQKEGGDSKECTTPQMQLNLALYTLNPLNIYRNQTTTSAQOHLT		
TRANSLATION OF LNCAP-POL-GENA-GOODA	(411)	QAIVERTNRTLKTQLVKQKEGGDSKECTTPQMQLNLALYTLNPLNIYRNQTTTSAQOHLT		
TRANSLATION OF ORF111-10	(417)	QAIVERTNRTLKTQLVKQKEGGDSKECTTPQMQLNLALYTLNPLNIYRNQTTTSAQOHLT		
PGD-P1	(17)	-----		
PGD-P2	(17)	-----		
PGDP3	(1)	-----		
CONSENSUS	(661)	QAIVERTNRTLKTQLVKQKEGGDSKECTTPQMQLNLALYTLNPLNIYRNQTTTSA QHLT		

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FIGURE 8 CONTD...

GI_4185939_EMB_CAA76879.1	(718)	GKKNSPHEGKLIWWDKSNKNTWEIGKVITWGRGFACVSPGENQLPVWIPTRLHKFYNEPI	780
GI_4185943_EMB_CAA76882.1	(721)	GKKNSPHEGKLIWWDKSNKNTWEIGKVITWGRGFACVSPGENQLPVWIPTRLHKFYNEPI	
GI_4185947_EMB_CAA76885.1	(718)	GKKNSPHEGKLIWWDKSNKNTWEIGKVITWGRGFACVSPGENQLPVWIPTRLHKFYNEPI	
GI_5931705_EMB_CAB56603.1	(691)	GKKNSPHEGKLI	
ENV OF AB047240	(424)	GKKHSPHEGKLIWWDKSNKNTWEIGKVITWGRGFACVSPGENQLPVWIPTRLHKFYNEPI	
TRANSLATION OF P386TOP-LINK	(31)	-----	
TRANSLATION OF POL349-LINK	(28)	-----	
LNCAP-GENOMEA-POLORF	(471)	GKKHSPHEGKLIWWDKSNKNTWEIGKVITWGRGFACVSPGENQLPVWIPTRLHKFYNEPI	
TRANSLATION OF LNCAP-POL-GENA-GOODA	(471)	GKKHSPHEGKLIWWDKSNKNTWEIGKVITWGRGFACVSPGENQLPVWIPTRLHKFYNEPI	
TRANSLATION OF ORF111-10	(477)	GKKHSPHEGKLIWWDKSNKNTWEIGKVITWGRGFACVSPGENQLPVWIPTRLHKFYNEPI	
PGD-P1	(17)	-----	
PGD-P2	(17)	-----	
PGDP3	(4)	GKKNSPHEGKLIC	
CONSENSUS	(721)	GKK SPHEGKLIWWDK SNKNTWEIGKVITWGRGFACVSPGENQLPVWIPTRLHKFYNEPI	
GI_4185939_EMB_CAA76879.1			781
GI_4185943_EMB_CAA76882.1	(778)	RDAAKSTSAETETS	840
GI_4185947_EMB_CAA76885.1	(781)	GDAKSTSAETETP	
GI_5931705_EMB_CAB56603.1	(778)	RDAAKSTSAETETS	
ENV OF AB047240	(703)	-----	
TRANSLATION OF P386TOP-LINK	(484)	GDAKRASTEMVTPVTWMDNPIEVYVNDVSVVPGPTDDRCPAKPEEGBMMINISIVYRYP	
TRANSLATION OF POL349-LINK	(31)	-----	
LNCAP-GENOMEA-POLORF	(28)	-----	
TRANSLATION OF LNCAP-POL-GENA-GOODA	(531)	GDAKRASTEMVTPVTWMDNPIEVYVNDVSVVPGPTDDRCPAKPEEGBMMINISIVYRYP	
TRANSLATION OF ORF111-10	(531)	GDAKRASTEMVTPVTWMDNPIEVYVNDVSVVPGPTDDRCPAKPEEGBMMINISIVYRYP	
PGD-P1	(537)	GDAKRASTEMVTPVTWMDNPIEVYVNDVSVVPGPTDDRCPAKPEEGBMMINISIVYRYP	
PGD-P2	(17)	-----	
PGDP3	(17)	-----	
CONSENSUS	(17)	-----	
GI_4185939_EMB_CAA76879.1	(781)	DAKK S E T	
GI_4185943_EMB_CAA76882.1			841
GI_4185947_EMB_CAA76885.1	(792)	-----	900
GI_5931705_EMB_CAB56603.1	(795)	-----	
ENV OF AB047240	(792)	-----	
TRANSLATION OF P386TOP-LINK	(703)	-----	
TRANSLATION OF POL349-LINK	(544)	PICLGRAPGCLMPAVQNWLVVEVPTVSPNSRFTYHVMVSGMSLRPRVNYLQDFSYQRLKFR	
LNCAP-GENOMEA-POLORF	(31)	-----	
TRANSLATION OF LNCAP-POL-GENA-GOODA	(28)	-----	
TRANSLATION OF ORF111-10	(591)	PICLGRAPGCLMPAVQNWLVVEVPTVSPNSRFTYHVMVSGMSLRPRVNYLQDFSYQRLKFR	
PGD-P1	(591)	PICLGRAPGCLMPAVQNWLVVEVPTVSPNSRFTYHVMVSGMSLRPRVNYLQDFSYQRLKFR	
PGD-P2	(597)	PICLGRAPGCLMPAVQNWLVVEVPTVSPNSRFTYHVMVSGMSLRPRVNYLQDFSYQRLKFR	
PGDP3	(17)	-----	
CONSENSUS	(17)	-----	
GI_4185939_EMB_CAA76879.1	(17)	-----	
GI_4185943_EMB_CAA76882.1	(17)	-----	
GI_4185947_EMB_CAA76885.1	(17)	-----	
GI_5931705_EMB_CAB56603.1	(17)	-----	
ENV OF AB047240	(604)	PKGKPCPKBI PKESKNTVLVWEECVANSVILQNNPEGTIDWAPRGQFYHNCSSGQTQS	
TRANSLATION OF P386TOP-LINK	(31)	-----	
TRANSLATION OF POL349-LINK	(28)	-----	
LNCAP-GENOMEA-POLORF	(651)	PKGKPCPKBI PKESKNTVLVWEECVANSVILQNNPEGTIDWAPRGQFYHNCSSGQTQS	
TRANSLATION OF LNCAP-POL-GENA-GOODA	(651)	PKGKPCPKBI PKESKNTVLVWEECVANSVILQNNPEGTIDWAPRGQFYHNCSSGQTQS	
TRANSLATION OF ORF111-10	(657)	PKGKPCPKBI PKESKNTVLVWEECVANSVILQNNPEGTIDWAPRGQFYHNCSSGQTQS	
PGD-P1	(17)	-----	
PGD-P2	(17)	-----	
PGDP3	(17)	-----	
CONSENSUS	(17)	-----	
GI_4185939_EMB_CAA76879.1	(901)	-----	960
GI_4185943_EMB_CAA76882.1	(792)	-----	961
GI_4185947_EMB_CAA76885.1	(795)	-----	962
GI_5931705_EMB_CAB56603.1	(792)	-----	963
ENV OF AB047240	(703)	-----	964
TRANSLATION OF P386TOP-LINK	(604)	PKGKPCPKBI PKESKNTVLVWEECVANSVILQNNPEGTIDWAPRGQFYHNCSSGQTQS	
TRANSLATION OF POL349-LINK	(31)	-----	
LNCAP-GENOMEA-POLORF	(28)	-----	
TRANSLATION OF LNCAP-POL-GENA-GOODA	(651)	PKGKPCPKBI PKESKNTVLVWEECVANSVILQNNPEGTIDWAPRGQFYHNCSSGQTQS	
TRANSLATION OF ORF111-10	(651)	PKGKPCPKBI PKESKNTVLVWEECVANSVILQNNPEGTIDWAPRGQFYHNCSSGQTQS	
PGD-P1	(657)	PKGKPCPKBI PKESKNTVLVWEECVANSVILQNNPEGTIDWAPRGQFYHNCSSGQTQS	
PGD-P2	(17)	-----	
PGDP3	(17)	-----	
CONSENSUS	(17)	-----	
GI_4185939_EMB_CAA76879.1	(901)	-----	961
GI_4185943_EMB_CAA76882.1	(816)	QEGRAANLITTKADAVSYKISREHKGDTNPREACSDDDCINGGKSPYCRSSCS	1020
GI_4185947_EMB_CAA76885.1	(819)	QESRAADLITTKADAVSYKISREHKGDTNPREACSDDDCINGGKSPYCRSSCS	
GI_5931705_EMB_CAB56603.1	(816)	QEGRAANLITTKADAVSYKISREHKGDTNPREACSDDDCINGGKSPYCRSSCS	
ENV OF AB047240	(703)	-----	
TRANSLATION OF P386TOP-LINK	(664)	CPSAQVSPVDSLTESLDKHKHKKLQSFYPWNEGKGI STPRPEIISPVGPEHPELMR	
TRANSLATION OF POL349-LINK	(31)	-----	
LNCAP-GENOMEA-POLORF	(28)	-----	
TRANSLATION OF LNCAP-POL-GENA-GOODA	(711)	CPSAQVSPVDSLTESLDKHKHKKLQSFYPWNEGKGI STPRPEIISPVGPEHPELMR	
TRANSLATION OF ORF111-10	(711)	CPSAQVSPVDSLTESLDKHKHKKLQSFYPWNEGKGI STPRPEIISPVGPEHPELMR	
PGD-P1	(717)	CPSAQVSPVDSLTESLDKHKHKKLQSFYPWNEGKGI STPRPEIISPVGPEHPELMR	
PGD-P2	(17)	-----	
PGDP3	(17)	-----	
CONSENSUS	(17)	-----	
GI_4185939_EMB_CAA76879.1	(961)	A D K P EWG I SP S	

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FIGURE 8 *CONTD...*

	1021	1035
GI_4185939_EMB_CAA76879.1	(873)	-----
GI_4185943_EMB_CAA76882.1	(876)	-----
GI_4185947_EMB_CAA76885.1	(873)	-----
GI_5931705_EMB_CAB56603.1	(703)	-----
ENV OF AB047240	(724)	LWPDITLEFGLEIKL
TRANSLATION OF P386TOP-LINK	(31)	-----
TRANSLATION OF POL349-LINK	(28)	-----
LNCAP-GENOMEA-POLORF	(764)	-----
TRANSLATION OF LNCAP-POL-GENA-GOODA	(771)	LWPDITLEFGLEIKL
TRANSLATION OF ORF111-10	(777)	LWPDITLEFGLEIKL
PGD-P1	(17)	-----
PGD-P2	(17)	-----
PGDP3	(17)	-----
CONSENSUS	(1021)	-----

FIGURE 9

	1	60
GI_4185940_EMB_CAA76880.1	(1)	-----
GI_4185944_EMB_CAA76883.1	(1)	-----
GI_4185948_EMB_CAA76886.1	(1)	-----
GI_5931706_EMB_CAB56604.1	(1)	-----
ENV OF AB047240	(1)	MATLIGQGRRLRIITLCGNDPDKITVPFNKQQVRQAFISSGAWQIGLANFLGIIDNHYPKT
TRANSLATION OF E207TOP-LINK	(1)	-----
TRANSLATION OF ENV287-LINK	(1)	-----
TRANSLATION OF T20.22A-23	(1)	-----
PGD-E1	(1)	-----
PGD-E2	(1)	-----
PGD-E3	(1)	-----
CONSENSUS	(1)	-----

	61	120
GI_4185940_EMB_CAA76880.1	(1)	-----
GI_4185944_EMB_CAA76883.1	(1)	-----
GI_4185948_EMB_CAA76886.1	(1)	-----
GI_5931706_EMB_CAB56604.1	(1)	-----
ENV OF AB047240	(61)	KIFQFLKLTWILPKITRREPLENALTVFTDGSSNGKAAAYTGPKERVIKTPYQSAQRAEL
TRANSLATION OF E207TOP-LINK	(1)	-----
TRANSLATION OF ENV287-LINK	(1)	-----
TRANSLATION OF T20.22A-23	(1)	-----
PGD-E1	(1)	-----
PGD-E2	(1)	-----
PGD-E3	(1)	-----
CONSENSUS	(61)	-----

	121	180
GI_4185940_EMB_CAA76880.1	(1)	-----
GI_4185944_EMB_CAA76883.1	(1)	-----
GI_4185948_EMB_CAA76886.1	(1)	-----
GI_5931706_EMB_CAB56604.1	(1)	-----
ENV OF AB047240	(121)	VAVITVLQDFDQPINIISDSAYVVQATRDVETALIKYSTDDHLNQLFNLLQQTVRKRNF
TRANSLATION OF E207TOP-LINK	(1)	-----
TRANSLATION OF ENV287-LINK	(1)	-----
TRANSLATION OF T20.22A-23	(1)	-----
PGD-E1	(1)	-----
PGD-E2	(1)	-----
PGD-E3	(1)	-----
CONSENSUS	(121)	-----

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FIGURE 9 CONTD...

GI_4185940_EMB_CAA76880.1_	(1)	181	240
GI_4185944_EMB_CAA76883.1_	(1)	-----	-----
GI_4185948_EMB_CAA76886.1_	(1)	-----	-----
GI_5931706_EMB_CAB56604.1_	(1)	-----	-----
ENV OF AB047240	(181)	FYITHIRAHTNLPGLPTKANEQADLLVSSAFIKAQELLALTHVNAAGLKNKFPDVTWKQAK	
TRANSLATION OF E207TOP-LINK	(1)	-----	-----
TRANSLATION OF ENV287-LINK	(1)	-----	-----
TRANSLATION OF T20.22A-23	(1)	-----	-----
PGD-E1	(1)	-----	-----
PGD-E2	(1)	-----	-----
PGD-E3	(1)	-----	-----
CONSENSUS	(181)	-----	-----
GI_4185940_EMB_CAA76880.1_	(1)	241	300
GI_4185944_EMB_CAA76883.1_	(1)	-----	-----
GI_4185948_EMB_CAA76886.1_	(1)	-----	-----
GI_5931706_EMB_CAB56604.1_	(1)	-----	-----
ENV OF AB047240	(241)	DIVQHCQTCCQVLHLSTQEAGVNPRLCPNALWQMDGTHVPSFGRLSYVHVTVDTYSHPIW	
TRANSLATION OF E207TOP-LINK	(1)	-----	-----
TRANSLATION OF ENV287-LINK	(1)	-----	-----
TRANSLATION OF T20.22A-23	(1)	-----	-----
PGD-E1	(1)	-----	-----
PGD-E2	(1)	-----	-----
PGD-E3	(1)	-----	-----
CONSENSUS	(241)	-----	-----
GI_4185940_EMB_CAA76880.1_	(1)	301	360
GI_4185944_EMB_CAA76883.1_	(1)	-----	-----
GI_4185948_EMB_CAA76886.1_	(1)	-----	-----
GI_5931706_EMB_CAB56604.1_	(1)	-----	-----
ENV OF AB047240	(301)	ATCQTGESTSHVKKHLLSCFAVMGVPEKIKTDNGPGYCSKAPQKFLSQWKISHTTGIPYN	
TRANSLATION OF E207TOP-LINK	(1)	-----	-----
TRANSLATION OF ENV287-LINK	(1)	-----	-----
TRANSLATION OF T20.22A-23	(1)	-----	-----
PGD-E1	(1)	-----	-----
PGD-E2	(1)	-----	-----
PGD-E3	(1)	-----	-----
CONSENSUS	(301)	-----	-----
GI_4185940_EMB_CAA76880.1_	(1)	361	420
GI_4185944_EMB_CAA76883.1_	(1)	-----MQRKAPPPRRRRHRNRAPLTHKMNMVTSSEQMKL	
GI_4185948_EMB_CAA76886.1_	(1)	-----MQRKAPPPRRRRHRNRAPLTHKMNMVTSSEQMKL	
GI_5931706_EMB_CAB56604.1_	(1)	-----MQRKAPPPRRRRHRNRAPLTHKMNMVTSSEQMKL	
ENV OF AB047240	(361)	SGGQAIVERTNRTLKTQLVKQKEGGDSKECTTPQMQLNLALYTLNFLNIYRNQTTTSAKQ	
TRANSLATION OF E207TOP-LINK	(1)	-----	-----
TRANSLATION OF ENV287-LINK	(1)	-----	-----
TRANSLATION OF T20.22A-23	(1)	-----MNPSEMQRKAPPPRRRRHRNRAPLTHKMNMVTSSEQMKL	
PGD-E1	(1)	-----	-----
PGD-E2	(1)	-----	-----
PGD-E3	(1)	-----	-----
CONSENSUS	(361)	-----	-----
GI_4185940_EMB_CAA76880.1_	(35)	421	480
GI_4185944_EMB_CAA76883.1_	(35)	PSTKKAEPPTWAQLKKLTQLATKYLENTKVTQTPESMLLAALMIVSMVVSLEPMPAGAAAA	
GI_4185948_EMB_CAA76886.1_	(35)	PSTKKAEPPTWAQLKKLTQLATKYLENTKVTQTPESMLLAALMIVSMVVSLEPMPAGAAAA	
GI_5931706_EMB_CAB56604.1_	(1)	-----	-----
ENV OF AB047240	(421)	HLTGKKHSPHEGKLIWWKDNKNKTWEIGKVITWGRGFACVSPGENQLFPVWIPTRHLPYN	
TRANSLATION OF E207TOP-LINK	(1)	-----	-----
TRANSLATION OF ENV287-LINK	(1)	-----	-----
TRANSLATION OF T20.22A-23	(40)	PSTKKAEPPTWAQLKKLTQLATKYLENTKVTQTPESMLLAALMIVSMVVSLEPMPAGAAAA	
PGD-E1	(1)	-----	-----
PGD-E2	(1)	-----	-----
PGD-E3	(1)	-----	-----
CONSENSUS	(421)	-----	-----

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FIGURE 9 CONTD...

GI_4185940_EMB_CAA76880.1_ (95) NYTYWAYVPPFP-~~TT~~RAVTWMDNPTEVYVNDVSVWVPGPIDDRCPAKPEEEGMMINISIGY 540
 GI_4185944_EMB_CAA76883.1_ (95) NYTYWAYVPPFP-~~TT~~RAVTWMDNPTEVYVNDVSVWVPGPTDDHCPAKPEEEGMMINISIGY
 GI_4185948_EMB_CAA76886.1_ (95) NYTYWAYVPPFP-~~TT~~RAVTWMDNPTEVYVNDVSVWVPGPIDDRCPAKPEEEGMMINISIGY
 GI_5931706_EMB_CAB56604.1_ (1) -----~~TT~~TPVTWMDNPTEVYVNDVSVWVPGPTDDRCAPAKPEEEGMMINISIGY
 ENV OF AB047240 (481) EPIGDAKKRASTE~~TT~~TPVTWMDNPTEVYVNDVSVWVPGPTDDRCAPAKPEEEGMMINISIVY
 TRANSLATION OF E207TOP-LINK (1) -----
 TRANSLATION OF ENV287-LINK (1) -----
 TRANSLATION OF T20.22A-23 (100) NYTYWAYVPPFP-~~TT~~RAVTWMDNPTEVYVNDVSVWVPGPIDDRCPAKPEEEGMMINISIGY
 PGD-E1 (1) -----
 PGD-E2 (1) -----
 PGD-E3 (1) -----
 CONSENSUS (481) LI VTWMDNP EVYVNDVSVWVPGP DD CPAKPEEEGMMINISI Y

GI_4185940_EMB_CAA76880.1_ (154) HYPPICLGRAPGCLMPAVQNWLVEVPTVSPICRFTYHVMVSGMSLRPRVNYLQDFSYQSRSL 600
 GI_4185944_EMB_CAA76883.1_ (154) RYPPICLGRAPGCLMPAVQNWLVEVPTVSPISRFTYHVMVSGMSLRPRVNYLQDFSYQSRF
 GI_4185948_EMB_CAA76886.1_ (154) HYPPICLGRAPGCLMPAVQNWLVEVPTVSPICRFTYHVMVSGMSLRPRVNYLQDFSYQSRSL
 GI_5931706_EMB_CAB56604.1_ (48) HYPPICLGRAPGCLMPAVQNWLVEVPTVSPNSRFTYHVMVSGMSLRPRVNYLQDFSYQSRSL
 ENV OF AB047240 (541) RYPPICLGRAPGCLMPAVQNWLVEVPTVSPNSRFTYHVMVSGMSLRPRVNYLQDFSYQSRSL
 TRANSLATION OF E207TOP-LINK (1) -----PSYQSRSL
 TRANSLATION OF ENV287-LINK (1) -----
 TRANSLATION OF T20.22A-23 (159) HYPPICLGRAPGCLMPAVQNWLVEVPTVSPICRFTYHVMVSGMSLRPRVNYLQDFSYQSRSL
 PGD-E1 (1) -----
 PGD-E2 (1) -----
 PGD-E3 (1) -----
 CONSENSUS (541) YPPICLGRAPGCLMPAVQNWLVEVPTVSP RFTYHVMVSGMSLRPRVN LQDFSYQSRSL

GI_4185940_EMB_CAA76880.1_ (214) KFRPKGKPCPKKEIPKESKNTTEVLWVEECVANSVILQNNFEGTIIIDWAPRGQFYHNCSSGQ 660
 GI_4185944_EMB_CAA76883.1_ (214) KFRPKGKPCPKKEIPKESKNTTEVLWVEECVANSVILQNNFEGTIIIDWAPRGQFYHNCSSGQ
 GI_4185948_EMB_CAA76886.1_ (214) KFRPKGKPCPKKEIPKESKNTTEVLWVEECVANSVILQNNFEGTIIIDWAPRGQFYHNCSSGQ
 GI_5931706_EMB_CAB56604.1_ (108) KFRPKGKPCPKKEIPKESKNTTEVLWVEECVANSVILQNNFEGTIIIDWAPRGQFYHNCSSGQ
 ENV OF AB047240 (601) KFRPKGKPCPKKEIPKESKNTTEVLWVEECVANSVILQNNFEGTIIIDWAPRGQFYHNCSSGQ
 TRANSLATION OF E207TOP-LINK (8) KFRPKGKPCPKKEIPKESKNTTEVL-----
 TRANSLATION OF ENV287-LINK (1) -----
 TRANSLATION OF T20.22A-23 (219) KFRPKGKPCPKKEIPKESKNTTEVLWVEECVANSVILQNNFEGTIIIDWAPRGQFYHNCSSGQ
 PGD-E1 (1) --RPGKPCPKKEIPKESK-----
 PGD-E2 (1) -----
 PGD-E3 (1) -----
 CONSENSUS (601) KFRPKGKPCPKKEIPKESKNTTEVLWVEECVANS VILQNNFEGTIIIDWAPRGQFYHNCSSGQ

GI_4185940_EMB_CAA76880.1_ (274) TQSCQSAQVSPAVIDSLTESLDKHKHKKLQSFYPWEWGEKGISTPRPKIISPVSQGEHPE 720
 GI_4185944_EMB_CAA76883.1_ (274) TQSCPSAQVSPAVIDSLTESLDKHKHKKLQSFYPWEWGEKGISTPRPKIISPVSQGEHPE
 GI_4185948_EMB_CAA76886.1_ (274) TQSCPSAQVSPAVIDSLTESLDKHKHKKLQSFYPWEWGEKGISTPRPKIISPVSQGEHPE
 GI_5931706_EMB_CAB56604.1_ (168) TQSCPSAQVSPAVIDSLTESLDKHKHKKLQSFYLWEEWEEKGISTPRPKIISPVSQGEHPE
 ENV OF AB047240 (661) TQSCPSAQVSPAVIDSLTESLDKHKHKKLQSFYPWEWGEKGISTPRPETISPVSQGEHPE
 TRANSLATION OF E207TOP-LINK (31) -----
 TRANSLATION OF ENV287-LINK (1) -----SDLTESLDKHKHKKLQSFYPWEWGEKGI-----
 TRANSLATION OF T20.22A-23 (279) TQSCPSAQVSPAVIDSLTESLDKHKHKKLQSFYPWEWGEKGISTPRPKIISPVSQGEHPE
 PGD-E1 (17) -----
 PGD-E2 (1) -----
 PGD-E3 (1) -----
 CONSENSUS (661) TQSC SAQVSPAVIDSLTESLDKHKHKKLQSFYPWEWGEKGISTPRP IISPVSQGEHPE

GI_4185940_EMB_CAA76880.1_ (334) LWRLTVASHHIREWSGNQTLSTRDRKPFYITDLNSSITVPLQSCIKPPYMLVVGNIIVIKP 780
 GI_4185944_EMB_CAA76883.1_ (334) LWRLTVASHHIREWSGNQTLSTRDRKPFYITDLNSSITVPLQSCIKPPYMLVVGNIIVIKP
 GI_4185948_EMB_CAA76886.1_ (334) LWRLTVASHHIREWSGNQTLSTRDRKPFYITDLNSSITVPLQSCIKPPYMLVVGNIIVIKP
 GI_5931706_EMB_CAB56604.1_ (228) LWRLTVASHHIREWSGNQTLSTRYRKPFYITDLNSITVPLQSCIKPPYMLVVGNIIVIKP
 ENV OF AB047240 (721) LW-----R-----W-----P-----
 TRANSLATION OF E207TOP-LINK (31) -----
 TRANSLATION OF ENV287-LINK (29) -----
 TRANSLATION OF T20.22A-23 (339) LWRLTVASHHIREWSGNQTLSTRDRKPFYITDLNSSITVPLQSCIKPPYMLVVGNIIVIKP
 PGD-E1 (17) -----
 PGD-E2 (1) -----LNSITVPLQSCIKPC-----
 PGD-E3 (1) -----
 CONSENSUS (721) LW RI LNS LTVPLQSCVKP

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FIGURE 9 CONTD...

GI_4185940_EMB_CAA76880.1_	(394)	DSQTITCENCRLTLCIDSTFNQHRILLVRAREGVWIPVSMRDPWEASPSVHILTEVLKG	781	840
GI_4185944_EMB_CAA76883.1_	(394)	DSQTITCENCRLTLCIDSTFNQHRILLVRAREGVWIPVSMRDPWEASPSVHILTEVLKG		
GI_4185948_EMB_CAA76886.1_	(394)	DSQTITCENCRLTLCIDSTFNQHRILLVRAREGVWIPVSMRDPWEASPSVHILTEVLKG		
GI_5931706_EMB_CAB56604.1_	(288)	ASQTITCENCRLFTCIDSTFNQHRILLVRAREGMWIPVSTDRPWEASPSVHILTEILKG		
ENV OF AB047240	(727)	-----DUTLEEGLEIKL-----		
TRANSLATION OF E207TOP-LINK	(31)	-----		
TRANSLATION OF ENV287-LINK	(29)	-----		
TRANSLATION OF T20.22A-23	(399)	DSQTITCENCRLTLCIDSTFNQHRILLVRAREGVWIPVSMRDPWEASPSVHILTEVLKG		
PGD-E1	(17)	-----		
PGD-E2	(17)	-----		
PGD-E3	(1)	-----		
CONSENSUS	(781)	DST W I L		
GI_4185940_EMB_CAA76880.1_	(454)	VLNRSKRFIFTLIAVIMGLIAVTATAAVAGVALHSSVQSVNFVNDWQKNSTRLWNSQSSI	841	900
GI_4185944_EMB_CAA76883.1_	(454)	VLNRSKRFIFTLIAVIMGLIAVTATAAVAGVALHSSVQSVNFVNDWQKNSTRLWNSQSSI		
GI_4185948_EMB_CAA76886.1_	(454)	VLNRSKRFIFTLIAVIMGLIAVTATAAVAGVALHSSVQSVNFVNDWQKNSTRLWNSQSSI		
GI_5931706_EMB_CAB56604.1_	(348)	VLNRSKRFIFTLIAVIMGLIAVTATAAVAGVALHSSVQSVNFVNYWQKNSTRLWNSQSSI		
ENV OF AB047240	(739)	-----		
TRANSLATION OF E207TOP-LINK	(31)	-----		
TRANSLATION OF ENV287-LINK	(29)	-----		
TRANSLATION OF T20.22A-23	(459)	VLNRSKRFIFTLIAVIMGLIAVTATAAVAGVALHSSVQSVNFVNDWQKNSTRLWNSQSSI		
PGD-E1	(17)	-----		
PGD-E2	(17)	-----		
PGD-E3	(1)	-----		
CONSENSUS	(841)			
GI_4185940_EMB_CAA76880.1_	(514)	DQKLANQINDLRQTVIWMGDRLMSLEHRFQLQCDWNTSDFCITPQIYNESEHHWDMVRRH	901	960
GI_4185944_EMB_CAA76883.1_	(514)	DQKLANQINDLRQTVIWMGDRLMSLEHRFQLQCDWNTSDFCITPQIYNESEHHWDMVRRH		
GI_4185948_EMB_CAA76886.1_	(514)	DQKLANQINDLRQTVIWMGDRLMSLEHRFQLQCDWNTSDFCITPQIYNESEHHWDMVRRH		
GI_5931706_EMB_CAB56604.1_	(408)	DQKLANQINDLRQTVIWMGDRLMTLEHHFQLQCDWNTSDFCITPQIYNESEHHWDMVRRH		
ENV OF AB047240	(739)	-----		
TRANSLATION OF E207TOP-LINK	(31)	-----		
TRANSLATION OF ENV287-LINK	(29)	-----		
TRANSLATION OF T20.22A-23	(519)	DQKLANQINDLRQTVIWMGDRLMSLEHRFQLQCDWNTSDFCITPQIYNESEHHWDMVRRH		
PGD-E1	(17)	-----		
PGD-E2	(17)	-----		
PGD-E3	(1)	-----		
CONSENSUS	(901)			
GI_4185940_EMB_CAA76880.1_	(574)	LQGREDNLTLDISKLEQIFEASKAHLNLPVGTAEIAGVADGLANLNPVTWVKTIGSTTI	961	1020
GI_4185944_EMB_CAA76883.1_	(574)	LQGREDNLTLDISKLEQIFEASKAHLNLPVGTAEIAGVADGLANLNPVTWVKTIGSTTI		
GI_4185948_EMB_CAA76886.1_	(574)	LQGREDNLTLDISKLEQIFEASKAHLNLPVGTAEIAGVADGLANLNPVTWVKTIGSTTI		
GI_5931706_EMB_CAB56604.1_	(468)	LQGREDNLTLDISKLEQIFEASKAHLNLPVGTAEIAGVADGLANLNPVTWIKTIRSTMI		
ENV OF AB047240	(739)	-----		
TRANSLATION OF E207TOP-LINK	(31)	-----		
TRANSLATION OF ENV287-LINK	(29)	-----		
TRANSLATION OF T20.22A-23	(579)	LQGREDNLTLDISKLEQIFEASKAHLNLPVGTAEIAGVADGLANLNPVTWVKTIGSTTI		
PGD-E1	(17)	-----		
PGD-E2	(17)	-----		
PGD-E3	(1)	-----		
CONSENSUS	(961)			
GI_4185940_EMB_CAA76880.1_	(634)	INLILILVCLFCLLLVCRCTQQLRRSDHRRERAMMTMAVLSKRKGGNVGKSKRDQIVTVSV	1021	1081
GI_4185944_EMB_CAA76883.1_	(634)	INLILILVCLFCLLLVCRCTQQLRRSDHRRERAMMTMAVLSKRKGGNVGKSKRDQIVTVSV		
GI_4185948_EMB_CAA76886.1_	(634)	INLILILVCLFCLLLVCRCTQQLRRSDHRRERAMMTMAVLSKRKGGNVGKSKRDQIVTVSV		
GI_5931706_EMB_CAB56604.1_	(528)	INLILIVVCLFCLLLVCRCTQQLRRSDIENG-		
ENV OF AB047240	(739)	-----		
TRANSLATION OF E207TOP-LINK	(31)	-----		
TRANSLATION OF ENV287-LINK	(29)	-----		
TRANSLATION OF T20.22A-23	(639)	INLILILVCLFCLLLVCRCTQQLRRSDHRRERAMMTMAVLSKRKGGNVGKSKRDQIVTVSV		
PGD-E1	(17)	-----		
PGD-E2	(17)	-----		
PGD-E3	(1)	-----RCTQQLRRSDHRRERA-----		
CONSENSUS	(1021)	RCTQQLRRSD		

FIGURE 9 CONTD...